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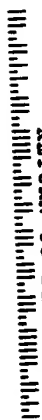
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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
09/903,188	07/11/2001	Edward M. De Robertis	510015-258	1059

7590 05/27/2004

Attention : Charles Berman
OPPENHEIMER WOLFF & DONNELLY
38th Floor
2029 Century Park East
Los Angeles, CA 90067-3024

EXAMINER

ROMEO, DAVID S

ART UNIT	PAPER NUMBER
----------	--------------

1647

DATE MAILED: 05/27/2004

Please find below and/or attached an Office communication concerning this application or proceeding.

Office Action Summary

Application No.

09/903,188

Applicant(s)

DE ROBERTIS ET AL

Examiner

David S Romeo

Art Unit

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-- The MAILING DATE of this communication appears on the cover sheet with the correspondence address --

Period for Reply

A SHORTENED STATUTORY PERIOD FOR REPLY IS SET TO EXPIRE 3 MONTH(S) FROM THE MAILING DATE OF THIS COMMUNICATION.

- Extensions of time may be available under the provisions of 37 CFR 1.136(a). In no event, however, may a reply be timely filed after SIX (6) MONTHS from the mailing date of this communication.
- If the period for reply specified above is less than thirty (30) days, a reply within the statutory minimum of thirty (30) days will be considered timely.
- If NO period for reply is specified above, the maximum statutory period will apply and will expire SIX (6) MONTHS from the mailing date of this communication.
- Failure to reply within the set or extended period for reply will, by statute, cause the application to become ABANDONED (35 U.S.C. § 133). Any reply received by the Office later than three months after the mailing date of this communication, even if timely filed, may reduce any earned patent term adjustment. See 37 CFR 1.704(b).

Status

- 1) ☒ Responsive to communication(s) filed on 22 August 2003.
- 2a) ☐ This action is **FINAL**. 2b) ☒ This action is non-final.
- 3) ☐ Since this application is in condition for allowance except for formal matters, prosecution as to the merits is closed in accordance with the practice under *Ex parte Quayle*, 1935 C.D. 11, 453 O.G. 213.

Disposition of Claims

- 4) ☒ Claim(s) 6-8, 11 and 12 is/are pending in the application.
- 4a) Of the above claim(s) 11 is/are withdrawn from consideration.
- 5) ☐ Claim(s) _____ is/are allowed.
- 6) ☒ Claim(s) 6-8 and 12 is/are rejected.
- 7) ☐ Claim(s) _____ is/are objected to.
- 8) ☒ Claim(s) 6-8, 11 and 12 are subject to restriction and/or election requirement.

Application Papers

- 9) ☐ The specification is objected to by the Examiner.
- 10) ☐ The drawing(s) filed on _____ is/are: a) ☐ accepted or b) ☐ objected to by the Examiner.
Applicant may not request that any objection to the drawing(s) be held in abeyance. See 37 CFR 1.85(a).
Replacement drawing sheet(s) including the correction is required if the drawing(s) is objected to. See 37 CFR 1.121(d).
- 11) ☐ The oath or declaration is objected to by the Examiner. Note the attached Office Action or form PTO-152.

Priority under 35 U.S.C. § 119

- 12) ☐ Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f).
- a) ☐ All b) ☐ Some * c) ☐ None of:
1. ☐ Certified copies of the priority documents have been received.
 2. ☐ Certified copies of the priority documents have been received in Application No. _____.
 3. ☐ Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)).

* See the attached detailed Office action for a list of the certified copies not received.

Attachment(s)

- 1) ☒ Notice of References Cited (PTO-892)
- 2) ☐ Notice of Draftsperson's Patent Drawing Review (PTO-948)
- 3) ☒ Information Disclosure Statement(s) (PTO-1449 or PTO/SB/08)
Paper No(s)/Mail Date 0402.
- 4) ☐ Interview Summary (PTO-413)
Paper No(s)/Mail Date. _____
- 5) ☐ Notice of Informal Patent Application (PTO-152)
- 6) ☐ Other: _____

DETAILED ACTION

The preliminary amendments filed 11/17/2003 and 07/11/2001 have been entered.

Claims 6-8, 11, 12 are pending.

5

Applicant's election of group I, claims 6-8, 12, in Paper No./the paper filed 08/22/2003 is acknowledged. Because applicant did not distinctly and specifically point out the supposed errors in the restriction requirement, the election has been treated as an election without traverse (MPEP § 818.03(a)).

10

Applicant's election of the polypeptide encoded by SEQ ID NO: 10 or comprising the amino acid sequence of SEQ ID NO: 9 species in Paper No./the paper filed 12/08/2003 is acknowledged.

15

Claim 11 is withdrawn from further consideration pursuant to 37 CFR 1.142(b) as being drawn to a nonelected invention, there being no allowable generic or linking claim. Election was made **without** traverse in Paper No./the paper filed 08/22/2003.

20 Claims 6-8, 12 are being examined. Claim 12 is being examined only to the extent that it reads upon the polypeptide encoded by SEQ ID NO: 10 or comprising the amino acid sequence of SEQ ID NO: 9 species.

Priority

Applicant has not complied with one or more conditions for receiving the benefit of an earlier filing date under 35 U.S.C. 120 as follows:

An application in which the benefits of an earlier application are desired must
5 contain a specific reference to the prior application(s) in the first sentence of the specification or in an application data sheet (37 CFR 1.78(a)(2) and (a)(5)). The specific reference to any prior nonprovisional application must include the relationship (i.e., continuation, divisional, or continuation-in-part) between the applications except when the reference is to a prior application of a CPA assigned the same application number.

10 It is acknowledged that the present application contains a specific reference to the 08/874,474 prior application in the first sentence of the specification. However, the specific reference to the 08/874,474 prior nonprovisional application does not include the relationship (i.e., continuation, divisional, or continuation-in-part) between the applications. The status of nonprovisional parent application(s) (whether patented or
15 abandoned) should also be included. If a parent application has become a patent, the expression "now Patent No. ____" should follow the filing date of the parent application. If a parent application has become abandoned, the expression "now abandoned" should follow the filing date of the parent application.

If a benefit claim to a provisional application is submitted without an indication
20 that an intermediate application directly claims the benefit of the provisional application and the instant nonprovisional application is not filed within the 12 month period or the relationship between each nonprovisional application is not indicated, the Office will not recognize such benefit claim and will not include the benefit claim on the filing receipt.

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Therefore, a petition under 37 CFR 1.78(a) and the surcharge set forth in 37 CFR 1.17(t) will be required if the intermediate application and the relationship of each nonprovisional application are not indicated within the period set forth in 37 CFR 1.78(a). Even if the Office has recognized a benefit claim by entering it into the Office's database and including it on applicant's filing receipt, the benefit claim is not a proper benefit claim under 35 U.S.C. 119(e) or 35 U.S.C. 120 and 37 CFR 1.78 unless the reference is included in an ADS or in the first sentence of the specification and all other requirements are met. Accordingly, the benefit of the filing dates of the 08/874,474 nonprovisional application and the 60/020,150 provisional application is denied.

It is acknowledged that Applicants submitted a petition on 11/17/2003 to accept an unintentionally delayed claim for priority. However, that petition has been dismissed. See the paper mailed 05/21/2004.

Claim Rejections - 35 USC § 102

The following is a quotation of the appropriate paragraphs of 35 U.S.C. 102 that form the basis for the rejections under this section made in this Office action:

A person shall be entitled to a patent unless –

(b) the invention was patented or described in a printed publication in this or a foreign country or in public use or on sale in this country, more than one year prior to the date of application for patent in the United States.

Claims 6-8, 12 are rejected under 35 U.S.C. 102(b) as being anticipated by De Robertis (N).

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This rejection is being made because the Office does not recognize Applicants benefit claims to the 08/874,474 nonprovisional application and the 60/020,150 provisional application, as discussed above.

De Robertis discloses a substantially pure protein characterized by a physiologically active form and comprising an amino acid sequence encoded by the DNA of SEQ ID NO: 10 (page 25, claim 6). De Robertis's SEQ ID NO: 10 is identical to the present application's SEQ ID NO: 10, as indicated below (Qy = the present application's SEQ ID NO: 10) (Db = De Robertis's SEQ ID NO: 10):

```

10  AAV14017
    ID  AAV14017 standard; cDNA; 1893 BP.
    XX
    AC  AAV14017;
    XX
    DT  09-JUL-1998 (first entry)
15  XX
    DE  Human "frazzled" frzb-1 cDNA.
    XX
    KW  Growth factor; frazzled; frzb-1; Wnts antagonist; human;
    KW  tumour suppressor; cancer; ds.
20  XX
    OS  Homo sapiens.
    XX
    FH  Key          Location/Qualifiers
    FT  CDS          61..1038
25  FT              /*tag= a
    FT              /product= frzb-1_protein
    XX
    PN  WO9748275-A1.
    XX
30  PD  24-DEC-1997.
    XX
    PF  19-JUN-1997;  97WO-US10942.
    XX
35  PR  18-JUN-1997;  97US-0878474.
    PR  20-JUN-1996;  96US-0020150.
    XX
    PA  (REGC ) UNIV CALIFORNIA.
    XX
40  PI  Bouwmeester T, De Robertis EM;
    XX
    DR  WPI; 1996-062760/06.
    DR  P-PSDB; AAW41254.
    XX
45  PT  New isolated growth factors - with neurotrophic, growth or
    PT  differentiation factor activity, tumour growth suppressor activity
    PT  or mesoderm differentiation activity
    XX
    PS  Claim 6; Fig 10; 48pp; English.
    XX
50  CC  The present sequence encodes the human growth factor protein
    CC  "frazzled" frzb-1. frzb-1 is an antagonist of Wnts in vivo, and
    CC  thus is believed to find utility as a tumour suppressor gene,
    CC  since overexpressed Wnt proteins cause cancer. Frzb-1 may also be a
55  CC  useful vehicle for solubilisation and therapeutic delivery of
    CC  complexed Wnt proteins.
    XX
    SQ  Sequence 1893 BP; 516 A; 438 C; 432 G; 507 T; 0 other;

60  Query Match          100.0%; Score 1893; DB 19; Length 1893;
    Best Local Similarity 100.0%; Pred. No. 0;
    Matches 1893; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

    Qy      1  GGCGGAGCGGGCCTTTTGGCGTCCACTGCGCGGCTGCACCTGCCCATCTGCCGGGATC 60
65  Db      1  GGCGGAGCGGGCCTTTTGGCGTCCACTGCGCGGCTGCACCTGCCCATCTGCCGGGATC 60

    Qy      61  ATGGTCTGCGCGCAGCCCGGAGGGATGCTGCTGCTGCGGCGCGGCTGCTTGCCTGGCT 120
    Db      61  ATGGTCTGCGCGCAGCCCGGAGGGATGCTGCTGCTGCGGCGCGGCTGCTTGCCTGGCT 120

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5 Db 61 ATGCTCTGCGGCAGCCCGGAGGGATGCTGCTGCTGCGGGCGGGCTGCTTGCCCTGGCT 120

Qy 121 GCTCTCTGCGCTGCTCGGGTGCCCGGGGCTCGGGCTGCAGCCTGTGAGCCCGTCCGCATC 180
|||||

10 Db 121 GCTCTCTGCGCTGCTCGGGTGCCCGGGGCTCGGGCTGCAGCCTGTGAGCCCGTCCGCATC 180
|||||

Qy 181 CCCCTGTGCAAGTCCCTGCCCTGGAACTGACTAAGATGCCAACCACTGCACCAAGC 240
|||||

15 Db 181 CCCCTGTGCAAGTCCCTGCCCTGGAACTGACTAAGATGCCAACCACTGCACCAAGC 240
|||||

Qy 241 ACTCAGGCCAAAGCCATCCTGGCCATCGAGCAGTTGAAAGTCTGCTGGGCACCCACTGC 300
|||||

20 Db 241 ACTCAGGCCAAAGCCATCCTGGCCATCGAGCAGTTGAAAGTCTGCTGGGCACCCACTGC 300
|||||

15 Qy 301 AGCCCGATCTGCTCTTCTCTCTGTCCTATGACGCGCCATCTGCACCATTGACTTC 360
|||||

20 Db 301 AGCCCGATCTGCTCTTCTCTCTGTCCTATGACGCGCCATCTGCACCATTGACTTC 360
|||||

20 Qy 361 CAGCAGAGCCCATCAAGCCCTGTAAGTCTGTGTGCGAGCGGGCCCGCAGGGCTGTGAG 420
|||||

25 Db 361 CAGCAGAGCCCATCAAGCCCTGTAAGTCTGTGTGCGAGCGGGCCCGCAGGGCTGTGAG 420
|||||

Qy 421 CCCATACTCATCAAGTACCGCCACTCGTGGCCGAGAACCTGGCCTGCGAGGAGCTGCCA 480
|||||

25 Db 421 CCCATACTCATCAAGTACCGCCACTCGTGGCCGAGAACCTGGCCTGCGAGGAGCTGCCA 480
|||||

Qy 481 GTGTACGACAGGGGCGTGTGCTCTCTCCGAGGCCATCGTTACTGCGGACGAGCTGAT 540
|||||

30 Db 481 GTGTACGACAGGGGCGTGTGCTCTCTCCGAGGCCATCGTTACTGCGGACGAGCTGAT 540
|||||

30 Qy 541 TTTCCTATGGATTCTAGTAACGAAACTGTAGAGGGGCAAGCAGTGAACGCTGTAAATGT 600
|||||

35 Db 541 TTTCCTATGGATTCTAGTAACGAAACTGTAGAGGGGCAAGCAGTGAACGCTGTAAATGT 600
|||||

35 Qy 601 AAGCCTATTAGAGCTACACAGAGACCTATTTCGGAAACAATTACAATATGTCTTCGG 660
|||||

40 Db 601 AAGCCTATTAGAGCTACACAGAGACCTATTTCGGAAACAATTACAATATGTCTTCGG 660
|||||

40 Qy 661 GCTAAAGTTAAAGAGATAAAGACTAAGTGCCATGATGTGACTGCACTAGTGGAGGTGAAG 720
|||||

45 Db 661 GCTAAAGTTAAAGAGATAAAGACTAAGTGCCATGATGTGACTGCACTAGTGGAGGTGAAG 720
|||||

45 Qy 721 GAGATTCTAAAGTCTCTCTGTTAAACATTCCACGGGACACTGTCAACCTCTATACCAGC 780
|||||

50 Db 721 GAGATTCTAAAGTCTCTCTGTTAAACATTCCACGGGACACTGTCAACCTCTATACCAGC 780
|||||

50 Qy 781 TCTGGCTGCTCTGCGCTCCACTTAATGTTAATGAGGAATATATCATCATGGGCTATGAA 840
|||||

55 Db 781 TCTGGCTGCTCTGCGCTCCACTTAATGTTAATGAGGAATATATCATCATGGGCTATGAA 840
|||||

55 Qy 841 GATGAGGAACGTTCCAGATTACTCTTGTGGAAGGCTCTATAGCTGAGAAGTGGAGGAT 900
|||||

60 Db 841 GATGAGGAACGTTCCAGATTACTCTTGTGGAAGGCTCTATAGCTGAGAAGTGGAGGAT 900
|||||

55 Qy 901 CGACTCGTAAAAAGTTAAGCGCTGGGATATGAAGCTTCGTATCTTGGACTCAGTAAA 960
|||||

60 Db 901 CGACTCGTAAAAAGTTAAGCGCTGGGATATGAAGCTTCGTATCTTGGACTCAGTAAA 960
|||||

60 Qy 961 AGTGATTCTAGCAATAGTATTCCACTCAGAGTCAGAAGTCTGGCAGGAACCGAACCC 1020
|||||

65 Db 961 AGTGATTCTAGCAATAGTATTCCACTCAGAGTCAGAAGTCTGGCAGGAACCGAACCC 1020
|||||

65 Qy 1021 CGGCAAGCAGCAACTAAATCCGAAATACAAAAAGTAACACAGTGGACTTCCTATTAAAG 1080
|||||

70 Db 1021 CGGCAAGCAGCAACTAAATCCGAAATACAAAAAGTAACACAGTGGACTTCCTATTAAAG 1080
|||||

70 Qy 1081 ACTTACTTGCACTGCTGGACTAGCAAGGAAATGCACTATTGCACATCATATTCTATT 1140
|||||

75 Db 1081 ACTTACTTGCACTGCTGGACTAGCAAGGAAATGCACTATTGCACATCATATTCTATT 1140
|||||

75 Qy 1141 GTTTACTATAAAAAATCATGTGATAACTGATTATTACTTCTGTTTCTCTTTGGTTTCTGC 1200
|||||

80 Db 1141 GTTTACTATAAAAAATCATGTGATAACTGATTATTACTTCTGTTTCTCTTTGGTTTCTGC 1200
|||||

75 Qy 1201 TTCTCTCTCTCTCAACCCCTTTGTAATGGTTTGGGGGCGAGCTCTAAGTATATTGTGA 1260
|||||

80 Db 1201 TTCTCTCTCTCTCAACCCCTTTGTAATGGTTTGGGGGCGAGCTCTAAGTATATTGTGA 1260
|||||

80 Qy 1261 GTTTTCTATTTCATAATCATGAGAAAACTGTTCTTTTGCAATAATAAATAAATAAACA 1320
|||||

85 Db 1261 GTTTTCTATTTCATAATCATGAGAAAACTGTTCTTTTGCAATAATAAATAAATAAACA 1320
|||||

85 Qy 1321 TGCTGTTACCAAGGCTCTTTGCTGAGTCTCCAGATGTTAATTACTTTCTGCACCCCAA 1380
|||||

90 Db 1321 TGCTGTTACCAAGGCTCTTTGCTGAGTCTCCAGATGTTAATTACTTTCTGCACCCCAA 1380
|||||

90 Qy 1381 TTGGGAATGCAATATTGGATGAAAAAGAGAGGTTTCTGGTATTACAGAAAGCTAGATATG 1440
|||||

95 Db 1381 TTGGGAATGCAATATTGGATGAAAAAGAGAGGTTTCTGGTATTACAGAAAGCTAGATATG 1440
|||||

90 Qy 1441 CCTTAAACATACTCTGCGATCTAATTACAGCCTTATTTTGTATGCCTTTGGGCATT 1500
|||||

95 Db 1441 CCTTAAACATACTCTGCGATCTAATTACAGCCTTATTTTGTATGCCTTTGGGCATT 1500
|||||

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5 QY 1501 CTCTCATGCTTAGAAAGTTCCAAATGTTTATAAAGGTAAAAATGGCAGTTTGAAAGTCAAA 1560
 Db 1501 CTCTCATGCTTAGAAAGTTCCAAATGTTTATAAAGGTAAAAATGGCAGTTTGAAAGTCAAA 1560

10 QY 1561 TGTCAATAGGCAAGCAATCAAGCACCAGGAAGTGTATGAGGAAACACCCCAAGA 1620
 Db 1561 TGTCAATAGGCAAGCAATCAAGCACCAGGAAGTGTATGAGGAAACACCCCAAGA 1620

15 QY 1621 TGAATTATTTTGGAGCTGTGAGGAAGTAAATAAATAGGAGCTTAAGAAAGAACATTTT 1680
 Db 1621 TGAATTATTTTGGAGCTGTGAGGAAGTAAATAAATAGGAGCTTAAGAAAGAACATTTT 1680

20 QY 1681 GCCTGATTGAGAAGCACAACCTGAAACCAAGTAGCCGCTGGGGTGTTAATGGTAGCATTCTT 1740
 Db 1681 GCCTGATTGAGAAGCACAACCTGAAACCAAGTAGCCGCTGGGGTGTTAATGGTAGCATTCTT 1740

25 QY 1741 CTTTGGCAATACATTGATTGTTTCATGAATATATTAAATCAGCATTAGAGAAATGAATT 1800
 Db 1741 CTTTGGCAATACATTGATTGTTTCATGAATATATTAAATCAGCATTAGAGAAATGAATT 1800

QY 1801 ATAACTAGACATCTGCTGTTATCACCATAGTTTGTGTTAATTGCTTCCTTTTAAATAAA 1860
 Db 1801 ATAACTAGACATCTGCTGTTATCACCATAGTTTGTGTTAATTGCTTCCTTTTAAATAAA 1860

QY 1861 CCCATTGGTGAAAGTCAAAAAAAAAAAAAAAAAA 1893
 Db 1861 CCCATTGGTGAAAGTCAAAAAAAAAAAAAAAAAA 1893.

30 De Robertis's SEQ ID NO: 10 encodes the amino acid sequence of SEQ ID NO: 9
 and SEQ ID NO: 9 is the amino acid sequence of human frzb-1 (page 6, lines 29-31;
 Figures 9 and 10). De Robertis's SEQ ID NO: 9 is identical to the present application's
 SEQ ID NO: 9, as indicated below (Qy = the present application's SEQ ID NO: 9) (Db =
 De Robertis's SEQ ID NO: 9):

35 AAW41254
 ID AAW41254 standard; protein; 325 AA.
 XX
 AC AAW41254;
 XX

40 DT 09-JUL-1998 (first entry)
 XX
 DE Human "frazzled" frzb-1.
 XX

45 KW Growth factor; frazzled; frzb-1; Wnts antagonist; human;
 KW tumour suppressor; cancer.
 XX
 OS Homo sapiens.
 XX

50 PN WO9748275-A1.
 XX
 PD 24-DEC-1997.
 XX
 PF 19-JUN-1997; 97WO-US010942.
 XX

55 PR 20-JUN-1996; 96US-0020150P.
 PR 18-JUN-1997; 97US-00878474.
 XX
 PA (REGC) UNIV CALIFORNIA.
 XX

60 PI De Robertis EM, Bouwmeester T;
 XX
 DR WPI; 1998-062760/06.
 DR N-PSDB; AAV14017.
 XX

65 PT New isolated growth factors - with neurotrophic, growth or
 PT differentiation factor activity, tumour growth suppressor activity or
 PT mesoderm differentiation activity.
 XX

70 PS Claim 6; Fig 9; 48pp; English.
 XX
 CC The present sequence is the human growth factor protein "frazzled" frzb-
 CC 1. frzb-1 is an antagonist of Wnts in vivo, and thus is believed to find
 CC utility as a tumour suppressor gene, since overexpressed Wnt proteins
 CC cause cancer. Frzb-1 may also be a useful vehicle for solubilisation and

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CC therapeutic delivery of complexed Wnt proteins

XX

SQ Sequence 325 AA;

Query Match 100.0%; Score 1738; DB 2; Length 325;
 Best Local Similarity 100.0%; Pred. No. 7.1e-166;
 Matches 325; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 M VCGSPGGMLLLRAGLLAALCLLRVPGARAAACEPVRIPLCKSLPWNMTKMPNHLHHS 60
 Db 1 M VCGSPGGMLLLRAGLLAALCLLRVPGARAAACEPVRIPLCKSLPWNMTKMPNHLHHS 60
 QY 61 TQANAILAIEQFEGLLGTHCSPDLLFFLCAMYAPICTIDFQHEPIKPCKSVCKERARQGCR 120
 Db 61 TQANAILAIEQFEGLLGTHCSPDLLFFLCAMYAPICTIDFQHEPIKPCKSVCKERARQGCR 120
 QY 121 PILIKYRHSWPENLACEELPVYDRGVCISPEAIVTADGADFFMDSSNGNCRGASSERCKC 180
 Db 121 PILIKYRHSWPENLACEELPVYDRGVCISPEAIVTADGADFFMDSSNGNCRGASSERCKC 180
 QY 181 KP1RATQKTYFRNNYNYVIRAKVKEIKTKCHDVTAVVEVKEILKSSLVNI PRDTVNLYTS 240
 Db 181 KP1RATQKTYFRNNYNYVIRAKVKEIKTKCHDVTAVVEVKEILKSSLVNI PRDTVNLYTS 240
 QY 241 SGCLCPPLNVNEEYIIMGYEDEERSRLLLVEGSIAEKWKDRLGKKVKRWDMLRHLGLSK 300
 Db 241 SGCLCPPLNVNEEYIIMGYEDEERSRLLLVEGSIAEKWKDRLGKKVKRWDMLRHLGLSK 300
 QY 301 SDSSNSDSTQSQKSGRNSNPRQARN 325
 Db 301 SDSSNSDSTQSQKSGRNSNPRQARN 325.

De Robertis also discloses a complex comprising a substantially pure frzb-1 protein complexed with at least one Wnt protein (claim 12, page 26). Accordingly, De Robertis discloses a complex comprising a substantially pure frzb-1 protein comprising the amino acid sequence of SEQ ID NO: 9 complexed with at least one Wnt protein.

Claim Rejections - 35 USC § 112

The following is a quotation of the second paragraph of 35 U.S.C. 112:

The specification shall conclude with one or more claims particularly pointing out and distinctly claiming the subject matter which the applicant regards as his invention.

Claim 12 is rejected under 35 U.S.C. 112, second paragraph, as being indefinite for failing to particularly point out and distinctly claim the subject matter which applicant regards as the invention.

The present specification discloses that "substitutional, deletional, or insertional mutants of the novel polypeptides may be prepared by in vitro or recombinant methods and screened for immuno-crossreactivity with cerberus, frzb-1, or PAPC and for cerberus

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antagonist or agonist activity" (page 5, lines 31-35). Hence, it is unclear how to construe the term "frzb-1 protein" because it is unclear if "substitutional, deletional, or insertional mutants" are encompassed by the term "frzb-1 protein." The metes and bounds are not clearly set forth.

5

Conclusion

No claims are allowable.

ANY INQUIRY CONCERNING THIS COMMUNICATION OR EARLIER COMMUNICATIONS FROM THE EXAMINER SHOULD BE DIRECTED TO DAVID S. ROMEO WHOSE TELEPHONE NUMBER IS (571) 272-0890. THE EXAMINER CAN NORMALLY BE REACHED ON MONDAY THROUGH FRIDAY FROM 7:30 A.M. TO 4:00 P.M. IF ATTEMPTS TO REACH THE EXAMINER BY TELEPHONE ARE UNSUCCESSFUL, THE EXAMINER'S SUPERVISOR, GARY KUNZ, CAN BE REACHED ON (571) 272-0887.

IF SUBMITTING OFFICIAL CORRESPONDENCE BY FAX, APPLICANTS ARE ENCOURAGED TO SUBMIT OFFICIAL CORRESPONDENCE TO THE FOLLOWING TC 1600 BEFORE AND AFTER FINAL RIGHT FAX NUMBERS:

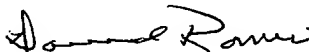
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CUSTOMERS ARE ALSO ADVISED TO USE CERTIFICATE OF FACSIMILE PROCEDURES WHEN SUBMITTING A REPLY TO A NON-FINAL OR FINAL OFFICE ACTION BY FACSIMILE (SEE 37 CFR 1.6 AND 1.8).

FAXED DRAFT OR INFORMAL COMMUNICATIONS SHOULD BE DIRECTED TO THE EXAMINER AT (571) 273-0890.

ANY INQUIRY OF A GENERAL NATURE OR RELATING TO THE STATUS OF THIS APPLICATION OR PROCEEDING SHOULD BE DIRECTED TO THE GROUP RECEPTIONIST WHOSE TELEPHONE NUMBER IS (703) 308-0196.



DAVID ROMEO
PRIMARY EXAMINER
ART UNIT 1647

DSR
MAY 26, 2004

Notice of References Cited

Application/Control No.

09/903,188

Applicant(s)/Patent Under
Reexamination
DE ROBERTIS ET AL

Examiner

David S Romeo

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U.S. PATENT DOCUMENTS

*		Document Number Country Code-Number-Kind Code	Date MM-YYYY	Name	Classification
	A	US-			
	B	US-			
	C	US-			
	D	US-			
	E	US-			
	F	US-			
	G	US-			
	H	US-			
	I	US-			
	J	US-			
	K	US-			
	L	US-			
	M	US-			

FOREIGN PATENT DOCUMENTS

*		Document Number Country Code-Number-Kind Code	Date MM-YYYY	Country	Name	Classification
	N	WO 97/48275	12-1997	WO	De Robertis et al.	----
	O					
	P					
	Q					
	R					
	S					
	T					

NON-PATENT DOCUMENTS

*		Include as applicable: Author, Title Date, Publisher, Edition or Volume, Pertinent Pages)
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	V	
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(54) Title: ENDODERM, CARDIAC AND NEURAL INDUCING FACTORS (57) Abstract Novel proteins have been designated "cerberus" and "frzb-1", respectively. Cerebus is expressed as a secreted peptide during embryogenesis of the <i>Xenopus</i> embryo, and is expressed specifically in the head organizer region. This new molecule has endodermal, cardiac, and neural tissue inducing activity, that should prove useful in therapeutic, diagnostic, and clinical applications requiring regeneration, differentiation, or repair of these and other tissues. Frzb-1 is a soluble antagonist of growth factors of the Wnt family that acts by binding to Wnt growth factors in the extracellular space. A third novel protein is termed PAPC which promotes the formation of dorsal mesoderm and somites in the embryo.		

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**ENDODERM, CARDIAC AND
NEURAL INDUCING FACTORS**

5 **Field of the Invention**

 The invention generally relates to growth factors, neurotrophic factors, and their inhibitors, and more particularly to several new growth factors with neural, endodermal, and cardiac tissue inducing activity, to complexes and compositions including the factors, and to DNA or RNA coding sequences for the factors. Further, one of the novel growth factors should be useful in tumor suppression gene therapy.

 This application claims the benefit of U.S. Provisional Application No. 60/020,150, filed June 20, 1996.

 This invention was made with Government support under grant contract number HD-21502, awarded by the National Institutes of Health. The Government has certain rights in this invention.

Background of the Invention

 Growth factors are substances, such as polypeptide hormones, which affect the growth of defined populations of animal cells *in vivo* or *in vitro*, but which are not nutrient substances. Proteins involved in the growth and differentiation of tissues may promote or inhibit growth, and promote or inhibit differentiation, and thus the general term "growth factor" includes cytokines, trophic factors, and their inhibitors.

Widespread neuronal cell death accompanies normal development of the central and peripheral nervous systems. Studies of peripheral target tissues during development have shown that neuronal cell death results from the competition among neurons for limiting amounts of survivor factors ("neurotrophic factors"). The earliest identified of these, nerve growth factor ("NGF"), is the most fully characterized and has been shown to be essential for the survival of sympathetic and neural crest-derived sensory neurons during early development of both chick and rat.

One family of neurotropic factors are the Wnts, which have dorsal axis-inducing activity. Most of the Wnt proteins are bound to cell surfaces. (See, e.g., Sokol et al., *Science*, 249, pp. 561-564, 1990.) Dorsal axis-inducing activity in *Xenopus* embryos by one member of this family (Xwnt-8) was described by Smith and Harland in 1991, *Cell*, 67, pp. 753-765. The authors described using RNA injections as a strategy for identifying endogenous RNAs involved in dorsal patterning to rescue dorsal development in embryos that were ventralized by UV irradiation.

Another member of the growth and neurotropic factor family was subsequently discovered and described by Harland and Smith, which they termed "noggin." (*Cell*, 70, pp. 829-840 (1992).) Noggin is a good candidate to function as a signaling molecule in Nieuwkoop's center, by virtue of its maternal transcripts, and in Spemann's organizer, through its zygotic organizer-specific expression. Besides noggin, other secreted factors may be involved in the organizer phenomenon.

Another *Xenopus* gene designated "chordin" that begins to be expressed in Spemann's organizer and that can completely rescue axial development in ventralized

embryos was described by Sasai et al., *Cell*, 79, pp. 779-790, 1994. In addition to dorsalizing mesoderm, chordin has the ability to induce neural tissue and its activities are antagonized by Bone Morphogenetic Protein-4 (Sasai et al., *Nature*, 376, pp. 333-336, 1995).

Therefore, the dorsal lip or Spemann's organizer of the *Xenopus* embryo is an ideal tissue for seeking novel growth and neurotrophic factors. New growth and neurotrophic factors are useful agents, particularly those that are secreted due to their ability to be used in physiologically active, soluble forms because these factors, their receptors, and DNA or RNA coding sequences therefore and fragments thereof are useful in a number of therapeutic, clinical, research, diagnostic, and drug design applications.

Summary of the Invention

In one aspect of the present invention, the sequence of the novel peptide that can be in substantially purified form is shown by SEQ ID NO:1. The *Xenopus* derived SEQ ID NO:1 has been designated "cerberus," and this peptide is capable of inducing endodermal, cardiac, and neural tissue development in vertebrates when expressed. The nucleotide sequence which, when expressed results in cerberus, is illustrated by SEQ ID NO:2. Since peptides of the invention induce endodermal, cardiac, and neural tissue differentiation in vertebrates, they should be able to be prepared in physiologically active form for a number of therapeutic, clinical, and diagnostic applications.

Cerberus was isolated during a search for molecules expressed specifically in Spemann's organizer containing a secretory signal sequence. In addition to cerberus, two other novel cDNAs were identified.

The *Xenopus* derived peptide that can be deduced from SEQ ID NO:3 encodes a novel protein we had earlier designated as "frazzled," a secreted protein of 318 amino acids that has dorsalizing activity in *Xenopus* embryos. We now designate the novel protein as "frzb-1." The gene for frzb-1 is expressed in many adult tissues of many animals, three of the cDNAs (*Xenopus*, mouse, and human) have been cloned by us. The accession numbers for the *Xenopus*, mouse, and human frzb-1 cDNA sequences of the gene now designated frzb-1 are U68059, U68058, and U68057, respectively. Frzb-1 has some degree of sequence similarity to the *Drosophila* gene frizzled which has been shown to encode a seven-transmembrane protein that can act both as a signalling and as a receptor protein (Vinson et al., *Nature*, 338, pp. 263-264, 1989; Vinson and Adler, *Nature*, 329, pp. 549-551, 1987). Vertebrate homologues of Frizzled have been isolated and they too were found to be anchored to the cell membrane by seven membrane spanning domains (Wang et al., *J. Biol. Chem.*, 271, pp. 4468-4476, 1996). Frzb-1 differs from the frizzled proteins in that it is an entirely soluble, diffusible secreted protein and therefore suitable as a therapeutic agent. The nucleotide sequence derived from *Xenopus* that, when expressed, results in frzb-1 protein is illustrated by SEQ ID NO:4. The frzb-1 protein derived from mouse is shown as SEQ ID NO:7, while the mouse frzb-1 nucleotide sequence is SEQ ID NO:8. The human derived frzb-1 protein is illustrated by SEQ ID NO:9, and the human frzb-1 nucleotide sequence is SEQ ID NO:10.

Frzb-1 is an antagonist of Wnts *in vivo*, and thus is believed to find utility as a tumor suppressor gene, since overexpressed Wnt proteins cause cancer. Frzb-1 may also be a useful vehicle for solubilization

and therapeutic delivery of Wnt proteins complexed with it.

The final cDNA isolated containing a signal sequence results in a peptide designated Paraxial
5 Protocadherin (PAPC). The cDNA for PAPC is a divergent member of the cadherin multigene family. PAPC is most related to protocadherin 43 reported by Sano et al., *The EMBO J.*, 12, pp. 2249-2256, 1993. As shown in SEQ ID NO:5, the PAPC gene encodes a transmembrane protein of
10 896 amino acids, of which 187 are part of an intracellular domain. PAPC is a cell adhesion molecule, and microinjection of PAPC mRNA constructs into *Xenopus* embryos suggest that PAPC acts as a molecule involved in mesoderm differentiation. A soluble form of the PAPC
15 extracellular domain is able to block muscle and mesoderm formation in *Xenopus* embryos. The nucleotide sequence encoding *Xenopus* PAPC is provided in SEQ ID NO:6.

Cerberus, frzb-1, or PAPC or fragments thereof
20 (which also may be synthesized by *in vitro* methods) may be fused (by recombinant expression or *in vitro* covalent methods) to an immunogenic polypeptide and this, in turn, may be used to immunize an animal in order to raise antibodies against the novel proteins. Antibodies
25 are recoverable from the serum of immunized animals. Alternatively, monoclonal antibodies may be prepared from cells from the immunized animal in conventional fashion. Immobilized antibodies are useful particularly in the diagnosis (*in vitro* or *in vivo*) or purification
30 of cerberus, frzb-1, or PAPC.

Substitutional, deletional, or insertional mutants of the novel polypeptides may be prepared by *in vitro* or recombinant methods and screened for immuno-crossreactivity with cerberus, frzb-1, or PAPC and for
35 cerberus antagonist or agonist activity.

Cerberus or frzb-1 also may be derivatized in vitro in order to prepare immobilized and labelled proteins, particularly for purposes of diagnosis of insufficiencies thereof, or for affinity purification of antibodies thereto.

Among applications for the novel proteins are tissue replacement therapy and, because frzb-1 is an antagonist of Wnt signaling, tumor suppression therapies. The cerberus receptor may define a novel signalling pathway. In addition, frzb-1 could permit the isolation of novel members of the Wnt family of growth factors.

Brief Description of the Drawings

Figure 1 illustrates the amino acid sequence (SEQ ID NO:1) of the Fig. 2 cDNA clone for cerberus;

Figure 2 illustrates a cDNA clone (SEQ ID NO:2) for cerberus derived from Xenopus. Sense strand is on top (5' to 3' direction) and the antisense strand on the bottom line (in the opposite direction);

Figures 3 and 4 show the amino acid and nucleotide sequence, respectively, of full-length frzb-1 from Xenopus (SEQ ID NOS:3 and 4);

Figures 5 and 6 show the amino acid and nucleotide sequence, respectively, of full-length PAPC from Xenopus (SEQ ID NOS:5 and 6);

Figures 7 and 8 show the amino acid and nucleotide sequence, respectively, of full-length frzb-1 from mouse (SEQ ID NOS:7 and 8); and

Figures 9 and 10 show the amino acid and nucleotide sequence, respectively, of full-length frzb-1 from human (SEQ ID NOS:9 and 10).

Detailed Description of the Preferred Embodiments

Among the several novel proteins and their nucleotide sequences described herein, is a novel endodermal, cardiac, and neural inducing factor in vertebrates that we have named "cerberus." When referring to cerberus, the present invention also contemplates the use of fragments, derivatives, agonists, or antagonists of cerberus molecules. Because cerberus has no homology to any reported growth factors, it is proposed to be the founding member of a novel family of growth factors with potent biological activities, which may be isolated using SEQ ID NO:2.

The amphibian organizer consists of several cell populations with region-specific inducing activities. On the basis of morphogenetic movements, three very different cell populations can be distinguished in the organizer. First, cells with crawling migration movements involute, fanning out to form the prechordal plate. Second, cells involute through the dorsal lip driven by convergence and extension movements, giving rise to the notochord of the trunk. Third, involution ceases and the continuation of mediolateral intercalation movements leads to posterior extension movements and to the formation of the tail notochord and of the chordoneural hinge. The three cell populations correspond to the head, trunk, and tail organizers, respectively.

The cerberus gene is expressed at the right time and place to participate in cell signalling by Spemann's organizer. Specifically, cerberus is expressed in the head organizing region that consists of crawling-migrating cells. The cerberus expressing region corresponds to the prospective foregut, including the liver and pancreas anlage, and the heart mesoderm.

Cerberus expression is activated by chordin, noggin, and organizer-specific homeobox genes.

Our studies were conducted in early embryos of the frog *Xenopus laevis*. The frog embryo is well suited to experiments, particularly experiments pertaining to generating and maintaining regional differences within the embryo for determining roles in tissue differentiation. It is easy to culture embryos with access to the embryos even at very early stages of development (preceding and during the formation of body pattern and differentiation) and the embryos are large. The initial work with noggin and chordin also had been in *Xenopus* embryos, and, as predicted, was highly conserved among vertebrates. Predictions based on work with *Xenopus* as to corresponding human noggin were proven true and the ability to clone the gene for human noggin was readily accomplished. (See the description of *Xenopus* work and cloning information in PCT application, published March 17, 1994, WO 9 405 800, and the subsequent human cloning based thereon in the PCT application, also published March 17, 1994, as WO 9 405 791.)

CLONING

The cloning of cerberus, frzb-1, and PAPC resulted from a comprehensive screen for cDNAs enriched in Spemann's organizer. Subtractive differential screening was performed as follows. In brief, poly A⁺ RNA was isolated from 300 dorsal lip and ventral marginal zone (VMZ) explants at stage 10½. After first strand cDNA synthesis approximately 70-80% of common sequences were removed by subtraction with biotinylated VMZ poly A⁺ RNA prepared from 1500 ventral gastrula halves. For differential screening, duplicate filters (2000 plaques per 15 cm plate, a total of 80,000 clones

screened) of an unamplified oriented dorsal lip library were hybridized with radiolabeled dorsal lip or VMZ cDNA. Putative organizer-specific clones were isolated, grouped by sequence analysis from the 5' end and whole-mount in situ hybridization, and subsequently classified into known and new dorsal-specific genes. Rescreening of the library (100,000 independent phages) with a cerberus probe resulted in the isolation of 45 additional clones, 31 of which had similar size as the longest one of the 11 original clones indicating that they were presumably full-length cDNAs. The longest cDNAs for cerberus, frzb-1, and PAPC were completely sequenced.

To explore the molecular complexity of Spemann's organizer we performed a comprehensive differential screen for dorsal-specific cDNAs. The method was designed to identify abundant cDNAs without bias as to their function. As shown in Table 1, five previously known cDNAs and five new ones were isolated, of which three (expressed as cerberus, frzb-1, and PAPC, respectively) had secretory signal sequences.

TABLE 1

	Previously Known Genes	Gene Product	No. of Isolates
	Chordin	novel secreted protein	70
	Goosecoid	homeobox gene	3
5	Pintallavis/XFKH-1	forkhead/transcription factor	2
	Xnot-2	homeobox gene	1
	Xlim-1	homeobox gene	1
	New Genes		
	Cerberus	novel secreted protein	11
10	PAPC	cadherin-like/transmembrane	2
	Frzb-1	novel secreted protein	1
	Sox-2	sry/transcription factor	1
	Fkh-like	forkhead/transcription factor	1

15 The most abundant dorsal-specific cDNA was
 chordin (chd), with 70 independent isolates. The second
 most abundant cDNA was isolated 11 times and named
 cerberus (after a mythological guardian dog with
 multiple heads). The cerberus cDNA encodes a putative
 20 secreted polypeptide of 270 amino acids, with an amino
 terminal hydrophobic signal sequence and a carboxy
 terminal cysteine-rich region (Fig. 1). Cerberus is
 expressed specifically in the head organizer region of
 the *Xenopus* embryo, including the future foregut.

25 An abundant mRNA found in the dorsal region of
 the *Xenopus* gastrula encodes the novel putative secreted
 protein we have designated as cerberus. Cerberus mRNA
 has potent inducing activity in *Xenopus* embryos, leading
 to the formation of ectopic heads. Unlike other
 organizer-specific factors, cerberus does not dorsalize
 30 mesoderm and is instead an inhibitor of trunk-tail
 mesoderm. Cerberus is expressed in the anterior-most

domain of the gastrula including the leading edge of the deep layer of the dorsal lip a region that, as shown here, gives rise to foregut and midgut endoderm. Cerberus promotes the formation of cement gland, olfactory placodes, cyclopic eyes, forebrain, and duplicated heart and liver (a foregut derivative). Because the pancreas is also derived from this foregut region, it is likely that cerberus induces pancreas in addition to liver. The expression pattern and inducing activities of cerberus suggest a role for a previously neglected region of the embryo, the prospective foregut endoderm, in the induction of the anterior head region of the embryo.

Turning to Fig. 1, *Xenopus cerberus* encodes a putative secreted protein transiently expressed during embryogenesis and the deduced amino acid sequence of *Xenopus cerberus* is shown. The signal peptide sequence and the nine cysteine residues in the carboxy-terminus are indicated in bold. Potential N-linked glycosylation sites are underlined. In database searches the cerberus protein showed limited similarity only to the mammalian Dan protein, a possible tumor suppressor proposed to be a DNA-binding protein.

Cerberus appears to be a pioneer protein, as its amino acid sequence and the spacing of its 9 cysteine residues were not significantly similar to other proteins in the databases (NCBI-Gen Bank release 93.0). We conclude that the second most abundant dorsal-specific cDNA encodes a novel putative secreted factor, which should be the founding member of a novel family of growth factors active in cell differentiation.

Cerberus Demarcates an Anterior Organizer Domain. Cerberus mRNA is expressed at low levels in the unfertilized egg, and zygotic transcripts start accumulating at early gastrula. Expression continues

during gastrula and early neurula, rapidly declining during neurulation. Importantly, cerberus expression starts about one hour after that of chd, suggesting that cerberus could act downstream of the chd signal.

5 Whole-mount *in situ* hybridizations reveal that expression starts in the yolky endomesodermal cells located in the deep layer of the organizer. The cerberus domain includes the leading edge of the most anterior organizer cells and extends into the lateral
10 mesoderm. The leading edge gives rise to liver, pancreas, and foregut in its midline, and the more lateral region gives rise to heart mesoderm at later stages of development.

15 Fig. 2 sets out the sequence of a full length *Xenopus* cDNA for cerberus.

 This entirely new molecule has demonstrated physiological properties that should prove useful in therapeutic, diagnostic, and clinical applications that require regeneration, differentiation, or repair of
20 tissues, such wound repair, neuronal regenerational or transplantation, supplementation of heart muscle differentiation, differentiation of pancreas and liver, and other applications in which cell differentiation processes are to be induced.

25 The second, novel, secreted protein we have discovered is called "frzb-1," which was shown to be a secreted protein in *Xenopus* oocyte microinjection experiments. Thus it provides a natural soluble form of the related extracellular domains of *Drosophila* and
30 vertebrate frizzled proteins. We propose that the latter proteins could be converted into active soluble forms by introducing a stop codon before the first transmembrane domain. We have noted that the cysteine-rich region of frzb-1 and frizzled contains some overall
35 structural homology with Wnt proteins using the Profile

Search homology program (Gribskov, *Meth. Enzymol.*, 183, pp. 146-159, 1990). This had raised the interesting possibility that frzb-1 could interact directly with Wnt growth factors in the extracellular space. This was
5 because we had found that when microinjected into *Xenopus* embryos, frzb-1 constructs have moderate dorsalizing activity, leading to the formation of embryos with enlarged brain and head, and shortened trunk. Somatic muscle differentiation, which requires
10 Xwnt-8, was inhibited. In the case of frzb-1, an attractive hypothesis, suggested by the structural homologies, was that it may act as an inhibitor of Wnt-8, a growth factor that has ventralizing activity in the *Xenopus* embryo (Christian and Moon, *Genes Dev.*, 7,
15 pp. 13-28, 1993). We have shown that frzb-1 can interact with Xwnt-8 and Wnt-1, and it is expected that it could also interact with other members of the Wnt family of growth factors, of which at least 15 members exist in mammals. In addition, a possible interaction
20 with Wnts was suggested by the recent discovery that dishevelled, a gene acting downstream of wingless, has strong genetic interaction with frizzled mutants in *Drosophila* (Krasnow et al., *Development*, 121, pp. 4095-4102, 1995). This possibility has been explored in
25 depth (Leyns et al., *Cell*, 88, pp. 747-756, March 21, 1997), because a soluble antagonist of the Wnt family of proteins is expected to be of great therapeutic value. Examples 1 and 2 illustrate tests that show antagonism of Xwnt-8 by binding to frzb-1.

30 Vertebrate homologues of Frizzled have been isolated and they too are anchored to the cell membrane by seven membrane spanning domains (Wang et al., *J. Biol. Chem.*, 271, pp. 4468-4476, 1996). Frzb-1 differs from the frizzled proteins in that it is an
35 entirely soluble, diffusible secreted protein and

therefore suitable as a therapeutic agent. The nucleotide sequence that when expressed results in frzb-1 protein is illustrated by SEQ ID NO:4.

5 SEQ ID NO:4 corresponds to the *Xenopus* homolog, but by using it in BLAST searches (and by cloning mouse frzb-1) we had been able to assemble the sequence of the entire mature human frzb-1 protein, SEQ ID NO:9. Indeed, human frzb-1 is encoded in six expressed sequence tags (ESTs) available in Genebank.

10 The human frzb-1 sequence can be assembled by overlapping in the 5' to 3' direction the ESTs with the following accession numbers in Genebank: H18848, R63748, W38677, W44760, H38379, and N71244. No function had yet been assigned to these EST sequences, but we

15 believe and thus propose here that human frzb-1 will have similar functions in cell differentiation to those described above for *Xenopus* frzb-1. The nucleotide sequence of human frzb-1 is shown in SEQ ID NO:10. The mouse frzb-1 protein and nucleotide sequences are

20 provided by SEQ ID NOS:7 and 8, respectively.

In particular, we believe that frzb-1 will prove useful in gene therapy of human cancer cells. In this rapidly developing field, one approach is to introduce vectors expressing anti-sense sequences to

25 block expression of dominant oncogenes and growth factor receptors. Another approach is to produce episomal vectors that will replicate in human cells in a controlled fashion without transforming the cells. For an example of the latter (an episomal expression vector

30 system for human gene therapy), reference is made to U.S. Patent 5,624,820, issued April 29, 1997, inventor Cooper.

Gene therapy now includes uses of human tumor suppression genes. For example, U.S. Patent 5,491,064,

35 issued February 13, 1996, discloses a tumor suppression

gene localized on chromosome 11 and described as potentially useful for gene therapy in cancers deleted or altered in their expression of that gene. Frzb-1 maps to chromosome 2q31-33 and loss of one copy of the
5 2q31-33 and loss of one copy of the 2q arm has been observed with high incidence in lung carcinomas, colo-rectal carcinomas, and neuroblastomas, which has lead to the proposal that the 2q arm carries a tumor suppressor gene. We expect frzb to be a tumor
10 suppressor gene, and thus to be useful in tumor suppression applications.

A number of applications for cerberus and frzb-1 are suggested from their pharmacological (biological activity) properties.

15 For example, the cerberus and frzb-1 cDNAs should be useful as a diagnostic tool (such as through use of antibodies in assays for proteins in cell lines or use of oligonucleotides as primers in a PCR test to amplify those with sequence similarities to the
20 oligonucleotide primer, and to determine how much of the novel protein is present).

Cerberus, of course, might act upon its target cells via its own receptor. Cerberus, therefore, provides the key to isolate this receptor. Since many
25 receptors mutate to cellular oncogenes, the cerberus receptor should prove useful as a diagnostic probe for certain tumor types. Thus, when one views cerberus as ligand in complexes, then complexes in accordance with the invention include antibody bound to cerberus,
30 antibody bound to peptides derived from cerberus, cerberus bound to its receptor, or peptides derived from cerberus bound to its receptor or other factors. Mutant forms of cerberus, which are either more potent agonists or antagonists, are believed to be clinically useful.

Such complexes of cerberus and its binding protein partners will find uses in a number of applications.

Practice of this invention includes use of an oligonucleotide construct comprising a sequence coding
5 for cerberus or frzb-1 and for a promoter sequence operatively linked in a mammalian or a viral expression vector. Expression and cloning vectors contain a nucleotide sequence that enables the vector to replicate in one or more selected host cells. Generally, in
10 cloning vectors this sequence is one that enables the vector to replicate independently of the host chromosomes, and includes origins of replication or autonomously replicating sequences. The well-known plasmid pBR322 is suitable for most gram negative
15 bacteria, the 2 μ plasmid origin for yeast and various viral origins (SV40, polyoma, adenovirus, VSV or BPV) are useful for cloning vectors in mammalian cells.

Expression and cloning vectors should contain a selection gene, also termed a selectable marker.
20 Typically, this is a gene that encodes a protein necessary for the survival or growth of a host cell transformed with the vector. The presence of this gene ensures that any host cell which deletes the vector will not obtain an advantage in growth or reproduction over
25 transformed hosts. Typical selection genes encode proteins that (a) confer resistance to antibiotics or other toxins, e.g. ampicillin, neomycin, methotrexate or tetracycline, (b) complement auxotrophic deficiencies.

Examples of suitable selectable markers for
30 mammalian cells are dihydrofolate reductase (DHFR) or thymidine kinase. Such markers enable the identification of cells which were competent to take up the cerberus nucleic acid. The mammalian cell transformants are placed under selection pressure which only the
35 transformants are uniquely adapted to survive by virtue

of having taken up the marker. Selection pressure is imposed by culturing the transformants under conditions in which the concentration of selection agent in the medium is successively changed. Amplification is the process by which genes in greater demand for the production of a protein critical for growth are reiterated in tandem within the chromosomes of successive generations of recombinant cells. Increased quantities of cerberus or frzb-1 can therefor be synthesized from the amplified DNA.

For example, cells transformed with the DHFR selection gene are first identified by culturing all of the transformants in a culture medium which contains methotrexate (Mtx), a competitive antagonist of DHFR. An appropriate host cell in this case is the Chinese hamster ovary (CHO) cell line deficient in DHFR activity, prepared and propagated as described by Urlaub and Chasin, *Proc. Nat. Acad. Sci.*, 77, 4216 (1980). The transformed cells then are exposed to increased levels of Mtx. This leads to the synthesis of multiple copies of the DHFR gene and, concomitantly, multiple copies of other DNA comprising the expression vectors, such as the DNA encoding cerberus or frzb-1. Alternatively, host cells transformed by an expression vector comprising DNA sequences encoding cerberus or frzb-1 and aminoglycoside 3' phosphotransferase (APH) protein can be selected by cell growth in medium containing an aminoglycosidic antibiotic such as kanamycin or neomycin or G418. Because eukaryotic cells do not normally express an endogenous APH activity, genes encoding APH protein, commonly referred to as neo resistant genes, may be used as dominant selectable markers in a wide range of eukaryotic host cells, by which cells transformed by the vector can readily be identified.

Expression vectors, unlike cloning vectors, should contain a promoter which is recognized by the host organism and is operably linked to the cerberus nucleic acid. Promoters are untranslated sequences located upstream from the start codon of a structural gene (generally within about 100 to 1000 bp) that control the transcription and translation of nucleic acid under their control. They typically fall into two classes, inducible and constitutive. Inducible promoters are promoters that initiate increased levels of transcription from DNA under their control in response to some change in culture conditions, e.g. the presence or absence of a nutrient or a change in temperature. At this time a large number of promoters recognized by a variety of potential host cells are well known. These promoters can be operably linked to cerberus encoding DNA by removing them from their gene of origin by restriction enzyme digestion, followed by insertion 5' to the start codon for cerberus or frzb-1.

Nucleic acid is operably linked when it is placed into a functional relationship with another nucleic acid sequence. For example, DNA for a presequence or secretory leader is operably linked to DNA for a polypeptide if it is expressed as a preprotein which participates in the secretion of the polypeptide; a promoter or enhancer is operably linked to a coding sequence if it affects the transcription of the sequence; or a ribosome binding site is operably linked to a coding sequence if it is positioned so as to facilitate translation. Generally, operably linked means that the DNA sequences being linked are contiguous and, in the case of a secretory leader, contiguous and in reading phase. Linking is accomplished by ligation at convenient restriction sites. If such sites do not

exit then synthetic oligonucleotide adapters or linkers are used in accord with conventional practice.

Transcription of the protein-encoding DNA in mammalian host cells is controlled by promoters obtained from the genomes of viruses such as polyoma, cytomegalovirus, adenovirus, retroviruses, hepatitis-B virus, and most preferably Simian Virus 40 (SV40), or from heterologous mammalian promoters, e.g. the actin promoter. Of course, promoters from the host cell or related species also are useful herein.

Cerberus and frzb-1 are clearly useful as a component of culture media for use in culturing cells, such as endodermal, cardiac, and nerve cells, *in vitro*. We believe cerberus and frzb-1 will find uses as agents for enhancing the survival or inducing the growth of liver, pancreas, heart, and nerve cells, such as in tissue replacement therapy.

The final cDNA isolated containing a signal sequence results in a peptide designated Paraxial Protocadherin (PAPC). The cDNA for PAPC is a divergent member of the cadherin multigene family. PAPC is most related to protocadherin 43 reported by Sano et al., *The EMBO J.*, 12, pp. 2249-2256, 1993. As shown in SEQ ID NO:5, the PAPC gene encodes a transmembrane protein of 896 amino acids, of which 187 are part of an intracellular domain. PAPC is a cell adhesion molecule, and microinjection of PAPC mRNA constructs into *Xenopus* embryos suggest that PAPC acts in mesoderm differentiation. The nucleotide sequence encoding *Xenopus* PAPC is provided in SEQ ID NO:6.

Therapeutic formulations of the novel proteins may be prepared for storage by mixing the polypeptides having the desired degree of purity with optional physiologically acceptable carriers, excipients or stabilizers, in the form of lyophilized cake or aqueous

solutions. Acceptable carriers, excipients or stabilizers are nontoxic to recipients at the dosages and concentrations employed, and include buffers such as phosphate, citrate, and other organic acids; anti-oxidants including ascorbic acid; low molecular weight (less than about 10 residues) polypeptides; proteins, such as serum albumin, gelatin or immunoglobulins. Other components can include glycine, glutamine, asparagine, arginine, or lysine; monosaccharides, disaccharides, and other carbohydrates including glucose, mannose, or dextrans; chelating agents such as EDTA; sugar alcohols such as mannitol or sorbitol; salt-forming counterions such as sodium; and/or nonionic surfactants such as Tween, Pluronic or PEG.

Polyclonal antibodies to the novel proteins generally are raised in animals by multiple subcutaneous (sc) or intraperitoneal (ip) injections of cerberus or frzb-1 and an adjuvant. It may be useful to conjugate these proteins or a fragment containing the target amino acid sequence to a protein which is immunogenic in the species to be immunized, e.g., keyhole limpet hemocyanin, serum albumin, bovine thyroglobulin, or soybean trypsin inhibitor using a bifunctional or derivatizing agent, for example, maleimidobenzoyl sulfosuccinimide ester (conjugation through cysteine residues), N-hydroxysuccinimide (through lysine residues), glutaraldehyde, succinic anhydride, SOCl_2 , or $\text{R}^1\text{N} = \text{C} = \text{NR}$.

Animals can be immunized against the immunogenic conjugates or derivatives by combining 1 mg or 1 μg of conjugate (for rabbits or mice, respectively) with 3 volumes of Freund's complete adjuvant and injecting the solution intradermally in multiple sites. One month later the animals are boosted with 1/5 to 1/10 the original amount of conjugate in Freund's complete

adjuvant by subcutaneous injection at multiple sites. Seven to 14 days later animals are bled and the serum is assayed for anti-cerberus titer. Animals are boosted until the titer plateaus. Preferably, the animal is
5 boosted with the conjugate of the same cerberus or frzb-1 polypeptide, but conjugated to a different protein and/or through a different cross-linking agent. Conjugates also can be made in recombinant cell culture as protein fusions. Also, aggregating agents such as
10 alum are used to enhance the immune response.

Monoclonal antibodies are prepared by recovering spleen cells from immunized animals and immortalizing the cells in conventional fashion, e.g. by fusion with myeloma cells or by EB virus transformation
15 and screening for clones expressing the desired antibody.

Antibodies are useful in diagnostic assays for cerberus, frzb-1, or PAPC or their antibodies and to identify family members. In one embodiment of a
20 receptor binding assay, an antibody composition which binds to all of a selected plurality of members of the cerberus family is immobilized on an insoluble matrix, the test sample is contacted with the immobilized antibody composition in order to adsorb all cerberus
25 family members, and then the immobilized family members are contacted with a plurality of antibodies specific for each member, each of the antibodies being individually identifiable as specific for a predetermined family member, as by unique labels such as
30 discrete fluorophores or the like. By determining the presence and/or amount of each unique label, the relative proportion and amount of each family member can be determined.

The antibodies also are useful for the
35 affinity purification of the novel proteins from

recombinant cell culture or natural sources. Antibodies that do not detectably cross-react with other growth factors can be used to purify the proteins free from these other family members.

5

EXAMPLE 1**Frzb-1 Antagonizes Xwnt-8 Non-Cell Autonomously**

To test whether frzb-1 can antagonize secondary axes caused by Xwnt-8 after secretion by injected cells, an experimental design was used. Thus, frzb-1 mRNA was injected into each of the four animal blastomeres of eight-cell embryos, and subsequently, a single injection of Xwnt-8 mRNA was given to a vegetal-ventral blastomere at the 16-32 cell stage. In two independent experiments, we found that injection of frzb-1 alone (n=13) caused mild dorsalization with enlargement of the cement gland in all embryos and that injection of Xwnt-8 alone (n=53) lead to induction of complete secondary axes in 67% of the embryos. However, injection of frzb-1 into animal caps abolished the formation of complete axes induced by Xwnt-8 (n=27), leaving only a residual 14% of embryos with very weak secondary axes. The double-injected embryos retained the enlarged cement gland phenotype caused by injection of frzb-1 mRNA alone. Because both mRNAs encode secreted proteins and were microinjected into different cells, we conclude that the antagonistic effects of frzb-1 and Xwnt-8 took place in the extracellular space after these proteins were secreted.

EXAMPLE 2**Membrane-Anchored Wnt-1 Confers Frzb-1 Binding**

To investigate a possible interaction between frzb-1 and Wnts, the first step was to insert an HA epitope tag into a Xenopus frzb-1 construct driven by the CMV (cytomegalovirus) promoter. Frzbl-HA was tested in mRNA microinjection assays in Xenopus embryos and found to be biologically active. Conditioned medium from transiently transfected cells contained up to 10 μ g/ml of Frzbl-HA (quantitated on Western blots using an HA-tagged protein standard).

Transient transfection of 293 cells has been instrumental in demonstrating interactions between wingless and frizzled proteins. We therefore took advantage of constructs in which Wnt-1 was fused at the amino terminus of CD8, generating a transmembrane protein containing biologically active Wnt-1 exposed to the extracellular compartment. A Wnt1CD8 cDNA construct (a generous gift of Dr. H. Varmus, NIH) was subcloned into the pcDNA (Invitrogen) vector and transfected into 293 cells. After incubation with Frzbl-HA-conditioned medium (overnight at 37°C), intensely labeled cells were observed by immunofluorescence. As a negative control, a construct containing 120 amino acids of Xenopus chordin, an unrelated secreted protein was used. Transfection of this construct produced background binding of Frzbl-HA to the extracellular matrix, both uniform and punctate. Cotransfection of Wnt1CD8 with pcDNA-LacZ showed that transfected cells stained positively for Frzbl-HA and LacZ. Since Wnt1CD8 contains the entire CD8 molecule, a CD8 cDNA was used as an additional negative control. After transfection with LacZ and full-length CE8, Frzbl-HA failed to bind to the transfected cells. Although most of our experiments

were carried out at 37°C, Frzb1-HA-conditioned medium also stained Wnt1CD8-transfected cells after incubation at 4°C for 2 hours.

Attempts to biochemically quantitate the binding of Frzb-1 to Wnt1CD8-transfected cells were unsuccessful due to high background binding to control cultures, presumably due to binding to the extracellular matrix. Thus, we were unable to estimate a K_D for the affinity of the Frzb-1/Wnt-1 interaction. However, when serial dilutions of conditioned medium containing Frzb1-HA were performed (ranging from 2.5×10^{-7} to 1.25×10^{-10} M), staining of Wnt1CD8-transfected cells was found at all concentrations.

Although we have been unable to provide biochemical evidence for direct binding between Wnts and frzb-1, this cell biological assay indicates that Frzb1-HA can bind, directly or indirectly, to Wnt-1 on the cell membrane in the 10^{-10} M range.

It is to be understood that while the invention has been described above in conjunction with preferred specific embodiments, the description and examples are intended to illustrate and not limit the scope of the invention, which is defined by the scope of the appended claims.

It is Claimed:

1. A substantially pure protein characterized by a physiologically active form and comprising an amino acid sequence encoded by the DNA of SEQ ID NO:2.
2. The protein as in claim 1 having neurotrophic, growth or differentiation factor activity.
3. A composition comprising the protein of claim 1 and a physiologically acceptable carrier with which the peptide is admixed.
4. An oligonucleotide construct comprising a sequence coding for a protein and an expression vector operatively linked therewith, the protein having neurotrophic, growth or differentiation factor activity and being expressible from SEQ ID NO:2.
5. The construct as in claim 4 wherein the expression vector is a mammalian or viral expression vector.
6. A substantially pure protein characterized by a physiologically active form and comprising an amino acid sequence encoded by the DNA of SEQ ID NO:4, SEQ ID NO:8, or SEQ ID NO:10.
7. The protein as in claim 6 having neurotrophic, growth or differentiation factor activity.
8. A composition comprising the protein of claim 6 and a physiologically acceptable carrier with which the protein is admixed.

9. An oligonucleotide construct comprising a sequence coding for a protein and an expression vector operatively linked therewith, the protein being expressible from SEQ ID NO:4, SEQ ID NO:8 or SEQ ID NO:10.

10. The construct as in claim 9 wherein the protein is expressible in soluble form.

11. The construct as in claim 9 wherein the expression vector is a mammalian or viral expression vector.

12. A complex comprising a substantially pure frzb-1 protein complexed with at least one Wnt protein.

13. A substantially pure protein characterized by a physiologically active form and comprising an amino acid sequence encoded by the DNA of SEQ ID NO:6.

14. The protein as in claim 13 having mesoderm differentiation activity.

15. A composition comprising the protein of claim 13 and a physiologically acceptable carrier with which the protein is admixed.

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MLLNVLRICI	IVCLVNDGAG	KHSEGRERTK	TYSLNSRGYF	40
RKERGARRSK	ILLVNTKGLD	EPHIGHGDFG	LVAELFDSTR	80
THTNRKEPDM	NKVLFSTVA	HGNKSARRKA	YNGSRRNIFS	120
RRSFDKRNT	VTEKPGAKMF	WNNFLVKMNG	APQNTSHGSK	160
AQEIMKEACK	TLPFTQNIVH	ENCDRMVIQN	NLCFGKCISL	200
HVPNQDDR	TCSHCLPSKF	TLNHLTLNCT	GSKNVVKVVM	240
MVEECTCEAH	KSNFHQTAQF	NMDTSTTLHH		270

Figure 1

SUBSTITUTE SHEET (RULE 26)

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GAATTCOCAG CAAGTCGCTC AGAAACACTG CAGGGTCTAG ATATCAIACA ATGTTACTAA	60
CTTAAGGGTC GTTCAGCGAG TCTTTGTGAC GTCCAGATC TATAGTATGT TACAATGATT	
ATGTAICTCAG GATCTGTATT ATCGTCTGCC TTGTGAATGA TGGAGCAGGA AAACACTCAG	120
TACATGAGTC CTAGACATAA TAGCAGACGG AACACTTACT ACCTCGTCCT TTTGTGAGTC	
AAGGACGAGA AAGGACAAAA ACATATTCAC TTAACAGCAG AGGTTACTTC AGAAAAGAAA	180
TTCTGCTCT TCTCTGTTTT TGTATAAGTG AATTGTCTGTC TCCAATGAAG TCTTTTCTTT	
GAGGAGCAGC TAGGAGCAAG ATTCTGCTGG TGAATACTAA AGGTCTTGAT GAACCCACAC	240
CTCTCGTGC ATCTCGTTC TAAGACGACC ACTTATGATT TCAGAACTA CTGGGGGTG	
TTGGGCATGG TGATTTTCGC TTAGTAGCTG AACTATTTGA TTCCACCAGA ACACATACAA	300
AACCCGTACC ACTAAAAGCG AATCATCGAC TTGATAAACT AAGGTGGTCT TGTGTATGTT	
ACAGAAAAGA GCCAGACATG AACAAAGTCA AGCTTTTCTC AACAGTTGCC CATGGAACA	360
TGTCTTTCT CGGTCTGTAC TTGTTTCAGT TCGAAAAGAG TTGTCAACGG GTACCTTTGT	
AAAGTGCAAG AAGAAAAGCT TACAATGGTT CTAGAAGGAA TATTTTTOCT CGCCGTTCTT	420
TTTACGTTT TCTTTTTCGA ATGTTACCAA GATCTTCCTT ATAAAAAGGA GCGGCAAGAA	
TTGATAAAG AAATACAGAG GTTACTGAAA AGCCTGGTGC CAAGATGTTT TGAACAATT	480
AACATTTTC TTTATGTCTC CAATGACTTT TCGGAOCACG GTTCTACAAG ACCTTGTTAA	
TTTTGGTTAA AATGAATGGA GCCCCACAGA ATACAAGCCA TGGCAGTAAA GCACAGGAAA	540
AAAACCAATT TTACTTACCT CGGGGTGTCT TATGTTGGT ACCGTCAATT CGTGTCTTT	
TAATGAAAGA AGCTTGCAAA ACCTTGTTTT TCACTCAGAA TATTGTACAT GAAAACGTG	600
ATTACTTTCT TCGAACGTTT TGGAACAAAA AGTGAGTCTT ATAACATGTA CTTTGTACAC	
ACAGGATGGT GATACAGAAC AATCTGTGCT TTGGTAAATG CATCTCTCTC CATGTTCCAA	660
TGTCTTACCA CTATGTCTTG TTAGACACGA AACCATTTAC GTAGAGAGAG GTACAAGGT	
ATCAGCAAGA TCGACGAAAT ACTTGTTCOC ATTGCTTGCC GTCCAAATTT ACCCTGAACC	720
TAGTGTCTCT AGCTGCTTTA TGAACAAAGG TAACGAACGG CAGGTTTAAA TGGGACTTGG	
ACCTGAOGCT GAATTGTACT GGATCTAAGA ATGTAGTAAA GGTGTGTCATG ATGGTAGAGG	780
TGGACTGCGA CTTAACATGA CTTAGATTCT TACATCATT CCAACAGTAC TACCATCTCC	
AATGCACGTG TGAAGCTCAT AAGAGCAACT TCACCAAAAC TGCACAGTTT AACATGGATA	840
TTACGTGCAC ACTTOGAGTA TTCTCGTTGA AGGTGGTTTG ACGTGTCAAA TTGTACCTAT	
CATCTACTAC OCTGCACCAT TAAAGGACTG CCATACAGTA TGGAAAATGCC CTTTGTGTGG	900
GTAGATGATG GGACGTGGTA ATTTCTGAC GGTATGTCTT ACCTTTACGG GAAAACAACC	
AAATATTTGTT ACATACTATG CATCTAAAGC ATTATGTTGC CTTCTATTTC ATATAACCAC	960
TTATAAACAA TGTATGATAC GTAGATTTCG TAATACAACG GAAGATAAAG TATATTGGTG	
ATGGAATAAG GATTGTATGA ATTATAATTA ACAAATGGCA TTTTGTGTAA CATGCAAGAT	1020
TACCTTATTC CTAACATACT TAATATTAA TGTTACCGT AAAACACATT GTACGTTCTA	

Figure 2A

SUBSTITUTE SHEET (RULE 28)

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CTCTGTTCCA	TCAGTTGCAA	GATAAAAGGC	AATATTTGTT	TGACTTTTTT	TCTACAAAAT	1080
GAGACAAGGT	AGTCAACGTT	CTATTTTCCG	TTATAAACAA	ACTGAAAAAA	AGATGTTTTA	
GAATACCCAA	ATATATGATA	AGATAATGGG	GTCAAAACTG	TTAAGGGGTA	ATGTAATAAT	1140
CTTATGGGTT	TATATACTAT	TCTATTACCC	CAGTTTTGAC	AATTCOCCAT	TACATTATTA	
AGGGACTAAG	TTTGCCCAGG	AGCAGTGACC	CATAACAACC	AATCAGCAGG	TATGATTTAC	1200
TCCCTGATTC	AAACGGGTCC	TCGTCACTGG	GTATTGTTGG	TTAGTCGTCC	ATACTAAATG	
TGGTCACCTG	TTTAAAAGCA	AACATCTTAT	TGGTTGCTAT	GGGTTACTGC	TTCTGGGCAA	1260
ACCAGTGGAC	AAATTTTCGT	TTGTAGAATA	ACCAACGATA	CCCAATGACG	AAGACCCGTT	
AATGTGTGCC	TCATAGGGGG	GTTAGTGTGT	TGTGTACTGA	ATAAATTGTA	TTTATTTTCAT	1320
TTACACACGG	AGTATCCCCC	CAATCACACA	ACACATGACT	TATTTAACAT	AAATAAAGTA	
TGTTACAAA	AAAAAAA					
ACAATGTTTT	TTTTTTTT					

Figure 2B

SUBSTITUTE SHEET (RULE 26)

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MSRTRKVDL LLLAIPGLAL LLLPNAYCAS CEPVRIPMCK SMPWNMTKMP NHLHHSTQAN	60
ALLAIEQFEG LLTTECSQDL LFFLCAMYAP ICTIDFQHEP IKPCKSV CER ARAGCEPILI	120
KYRHTWPESL ACEELPVYDR GVCISPEAIV TVEQGTDSMP DFSMDSNNGN CGSGREHCKC	180
KPMKATQKTY LKNNYNYVIR AKVKEVKVKC HDATAIVEVK EILKSSLVNI PKDTVTLYTN	240
SGCLCPQLVA NEEYIIMGYE DKERTRLLLV EGSIAEKWRD RLAKKVKRWD QKLRRPRKSK	300
DPVAPIPNKN SNSRQARS	

Figure 3

SUBSTITUTE SHEET (RULE 26)

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GAATTCOCTT TCACACAGGA CTCTGGCAG AGGTGAATGG TTAGCCCTAT GGATTGTT	60
CTTAAGGGAA AGTGTGTCT GAGGACCGTC TCCACTTACC AATCGGGATA CCTAAACCAA	
TGTTGATTTT GACACATGAT TGATTGCTTT CAGATAGGAT TGAAGGACTT GGATTTTAT	120
ACAACATAAA CTGTGTACTA ACTAAGGAAA GTCTATCCTA ACTTCCTGAA CCTAAAAATA	
CTAATCTGTC ACTTTTAAAT TATCTGAGTA ATTGTTTATT TTGTATTGGA TGGGACTAAA	180
GATTAGACG TGAAAAATTA ATAGACTCAT TAACAAGTAA AACATAACCT ACCCTGATTT	
GATAAACTTA ACTCCTTGCT TTTGACTTGC CCATAAATA TAAGGTGGGG TGAGTTGTAG	240
CTATTGAAT TGAGGAACGA AAACCTGAACG GGTATTTGAT ATTCCACCCC ACTCAACATC	
TTGCTTTTAC ATGTGCCAG ATTTTCCCTG TATTCCTGT ATTCCCTCTA AAGTAAGCCT	300
AACGAAATG TACACGGGTC TAAAAGGGAC ATAAGGGACA TAAGGGAGAT TTCATTGGA	
ACACATACAG GTTGGGCAGA ATAACAATGT CTGGAACAAG GAAAGTGGAC TCATTACTGC	360
TGTGTATGTC CAACCCGTCT TATTGTTACA GAGCTTGTTT CTTTCACCTG AGTAATGACG	
TACTGGCCAT ACCTGGACTG GCGCTTCTCT TATTACCAA TGCTTACTGT GCTTCGTGTG	420
ATGACCGTA TGGAACCTGAC CGCGAAGAGA ATAATGGGTT ACGAATGACA CGAAGCACAC	
AGCCTGTGCG GATCCCATG TGCAAACTTA TGCCATGGAA CATGACCAAG ATGCCCAACC	480
TOGGACACGC CTAGGGGTAC ACGTTTAGAT ACGGTACCTT GACTTGTTT TACGGGTGG	
ATCTOCACCA CAGCACTCAA GCCAATGCCA TCCTGGCAAT TGAACAGTTT GAAGGTTTGC	540
TAGAGGTGGT GTGTGAGTT CGGTTACGGT AGGAACGTTA ACTTGTCAA CTTCCAAACG	
TGACCACTGA ATGTAGOCAG GACCTTTTGT TCTTCTGTG TGCCATGTAT GCCCCATTT	600
ACTGGTGACT TACATCGGTC CTGGAAAACA AGAAGACAC ACGGTACATA CGGGGGTAAA	
GTACCATGTA TTTCAGCAT GAACCAATTA AGCCTTGCAA GTCCGTGTGC GAAAGGGCCA	660
CATGGTAGCT AAAGGTGTA CTGGTTAAT TOGGACGTT CAGGCACACG CTTTCCGGT	
GGGCGGCTG TGAGCCCAT CTCTAAAGT ACCGGCACAC TTGGCCAGAG AGCCTGGCAT	720
CCGGGCGAC ACTCGGGTAA GAGTATTCA TGGCCGTGTG AACCGGTCTC TOGGACGTA	
GTGAAGAGCT GCGGTATAT GACAGAGGAG TCTGCATCTC CCAGAGGCT ATCGTCACAG	780
CACCTCTCGA CGGGCATATA CTGTCTCTC AGACGTAGAG GGGTCTCGA TAGCAGTGT	
TGGAACAAGG AACAGATTCA ATGCCAGACT TCTCATGGA TTCAAACAAT GGAAATTGCG	840
ACCTTGTTCC TTGTCTAAGT TACGGTCTGA AGAGGTACCT AAGTTTGTTA CCTTTAAGC	
GAAGGGCAG GGAGCACTGT AAATGCAAGC CCATGAAGC AACCCAAAG ACGTATCTCA	900
CTTGGCGTC CCGGTGACA TTACGTTG GGTACTTCCG TTGGGTTTTT TGCATAGAT	
AGAATAATTA CAATTATGTA ATCAGAGCAA AAGTGAAGA GGTGAAGTG AAATGCCACG	960
TCTTATTAAT GTTAATACAT TAGTCTGTT TTCACTTCT CCACTTTCAC TTTACGGTGC	
ACGCAACAGC AATTGTGGAA GTAAAGGAGA TTCTCAAGTC TTCCCTAGTG AACATTCTA	1020
TGGTTGTG TTAACACCTT CATTTCTCT AAGAGTTCAG AAGGGATCAC TTGTAAGGAT	

Figure 4A

SUBSTITUTE SHEET (RULE 26)

AAGACACAGT GACACTGTAC ACCAACTCAG GCTGCTTGTG CCCCCAGCTT GTTGCCAATG	1080
TTCTGTGTCA CTGTGACATG TGGTTGAGTC CGACGAACAC GGGGGTCGAA CRAACGGTTAC	
AGGAATACAT AATTATGGGC TATGAAGACA AAGAGCGTAC CAGGCTTCTA CTAGTGGGAG	1140
TCCTTATGTA TTAATACCCG ATACTTCTGT TTCTCGCATG GTCOGAAGAT GATCACTTC	
GATCCTTGGC CGAAAAATGG AGAGATCGTC TTGCTAAGAA AGTCAAGCGC TGGGATCAAA	1200
CTAGGAACCG GCTTTTACC TCTCTAGCAG AACGATTCTT TCAGTTCGCG ACCCTAGTTT	
AGCTTCGACG TCCAGGAAA AGCAAAGACC CCGTGGCTCC AATTCCCAAC AAAACAGCA	1260
TCGAAGCTGC AGGGTCCTTT TCGTTTCTGG GGCACCGAGG TTAAGGGTTG TTTTGTCTG	
ATTCCAGACA AGCGCGTAGT TAGACTAACG GAAAGGTGTA TGGAACTCT ATGGACTTTG	1320
TAAGGTCTGT TCGCGCATCA ATCTGATTGC CTTCCACAT ACCTTTGAGA TACCTGAAAC	
AACTAAGAT TTGCATTGTT GGAAGAGCAA AAAAGAAATT GCACTACAGC ACGTTATATT	1380
TTTGATTCTA AACGTAACAA CTTTCTCGTT TTTTCTTTAA CGTGATGTGG TGCAATATAA	
CTATTGTTA CTACAAGAAG CTGGTTTAGT TGATTGTAGT TCTCCTTCC TTCTTTTTT	1440
GATAACAAAT GATGTTCTTC GACCAATCA ACTAACATCA AGAGGAAGG AAGAAAAAA	
TTATAACTAT AATTGCACGT GTTCCAGGC AATTGTTTTA TTCAACTTCC AGTGACAGAG	1500
AATATTGATA TAAAGTGCA CAAGGGTCCG TTAACAAAT AAGTTGAAGG TCACTGTCTC	
CAGTGA CTGA ATGTCTCAGC CTAAAGAAGC TCAATTCATT TCTGATCAAC TAATGGTGAC	1560
GTCAGTACT TACAGAGTCG GATTCTTCG AGTTAAGTAA AGACTAGTTG ATTACCACTG	
AAGTGTGGA TACTTGGGGA AAGTGAAC TAATGCAATGG TAAATCAGAG AAAAGTTGAC	1620
TTACAAACT ATGAACCCCT TTCACTTGAT TAACGTTACC ATTAGTCTC TTTTCAACTG	
CAATGTTGCT TTTCTGTAG ATGAACAAGT GAGAGATCAC ATTTAAATGA TGATCACTTT	1680
GTTACAACGA AAAGGACATC TACTTGTTCA CTCTCTAGTG TAAATTTACT ACTAGTGAA	
CCATTTAATA CTTTCAGCAG TTTAGTTAG ATGACATGTA GGATGCACCT AAATCTAAAT	1740
GGTAAATTAT GAAAGTCGTC AAAATCAATC TACTGTACAT CCTACGTGGA TTTAGATTTA	
ATTTTATCAT AAATGAAGAG CTGGTTTGA CTGTATGGTC ACTGTTGGGA AGGTAAATGC	1800
TAAATAGTA TTTACTTCTC GACCAATCT GACATACCAG TGACAACTT TCCATTTACG	
CTACTTTGTC AATTCTGTTT TAAAAATTGC CTAAATAAAT ATTAAGTCCT AAATAAAAAA	1860
GATGAACAG TTAAGACAAA ATTTTAAACG GATTTATTZA TAATTCAGGA TTTATTTTTT	
AAAAAAAAA AAAAA	
TTTTTTTTT TTTT	

Figure 4B
SUBSTITUTE SHEET (RULE 26)

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MLLLFRAIPM LLGLMVLQT DCEIAQYYID EEEPPGTVIA VLSQHSIFNT TDIPATNFRL	60
MKQFNNSLIG VRES DGQLSI MERIDREQIC RQSLHCNLAL DVVSFSKGF KLLNVKVEVR	120
DINDHSPHFP SEIMHVEVSE SSSVGTRIP L EIAIDEDVGS NSIQNFQISN NSHPSIDVLT	180
RADGVKYADL VLMRELDREI QPTYIMELLA MDGGVPSLSG TAVVNIRVLD FNDNSPVFER	240
STIAVDLVED APLGYLLLEL HATDDDEGVN GEIVYGFSTL ASQEVRLFK INSRTGSVTL	300
EGQVDFETKQ TYEFVQAQD LGPNELTATC KVTVHILDVN DNTPAITITP LTTVNAGVAY	360
IPETATKENF IALISTTDRA SGSNGQVRCT LYGHEHFKLQ QAYEDSYMIV TTSTLDRENI	420
AAYSLTVVAE DLGFP SLKTK KYITVKVSE DNDAPVFSKP QYEASILENN APGSYITTVI	480
ARDSDSQNG KVNTRLVDK VMGQSLTFV SLDADSGVLR AVRSIDYEKL KQDFEIEAA	540
DNGIPQLSTR VQIANLRIVDQ NDNCPVITNP LLNNGSGEVL LPISAPQNYL VFQLKAEDSD	600
EGHNSQLFYT ILRDP SRLFA INKESGEVFL KKQLNSDHSE DLSIVVAVYD LGRPSLSTNA	660
TVKFILTDSF PSNVEVVILQ PSAEEQHQID MSIIFIAVLA GGCALLLLAI FVVACTCKKK	720
AGEFKQVPEQ HGTCEERLL STPSQSVSS SLSQSESCQL SINTESENC S VSSNQEQHQ	780
TGIKHSISVP SYHTSGWHL D NCAMSI SGHS HMGHISTKVQ WAKEIVTSM T VTLILVENQK	840
RRALSSQCRH KPVLTQMNQ QGSDMPITIS ATESTRVQKM GTACNMKRA IDCLTL	

Figure 5
SUBSTITUTE SHEET (RULE 26)

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GAATTCOCAG	AGATGAACTC	CTTGAGATTG	TTTAAATGA	CTGCAGGTCT	GGAAGGATTC	60
CTTAAGGGTC	TCTACTTGAG	GAACCTAAC	AAAATTTACT	GACGTCCAGA	CCTTCCTAAG	
ACATTGCCAC	ACTGTTTCTA	GGCATGAAAA	AACGTCAAGT	TTCAACTTTG	TTTTTGGTGC	120
TGTAACGGTG	TGACAAAGAT	CCGTACTTTT	TTGACGTTCA	AAGTTGAAAC	AAAAACCAAG	
AACTTTGATT	CTTCAAGATG	CTGCTTCTCT	TCAGAGCCAT	TCCAATGCTG	CTGTTGGGAC	180
TTGAAACTAA	GAAGTTCTAC	GACGAAGAGA	AGTCTCGGTA	AGGTTACGAC	GACAACCCCTG	
TGATGGTTTT	ACAAACAGAC	TGTGAAATTG	CCAGTACTA	CATAGATGAA	GAAGAACCCC	240
ACTACCAAAA	TGTTTGTCTG	ACACTTTAAC	GGGTCATGAT	GTATCTACTT	CTTCTTGGGG	
CTGGCACTGT	AATTGCAGTG	TTGTCACAAC	ACTCCATATT	TAACACTACA	GATATACCTG	300
GACCGTGACA	TTAACGTCAC	AACAGTGTG	TGAGGTATAA	ATTGTGATGT	CTATATGGAC	
CAACCAATTT	CCGTCTAATG	AAGCAATTTA	ATAATTOCCT	TATCGGAGTC	CGTGAGAGTG	360
GTTGGTTAAA	GGCAGATTAC	TTCTGTAAAT	TATTAAGGGA	ATAGCCCTCAG	GCACTCTCAC	
ATGGGCAGCT	GAGCATCATG	GAGAGGATTG	ACCGGGAGCA	AATCTGCAGG	CAGTCCCTTC	420
TACCCGTGGA	CTCGTAGTAC	CTCTCCTAAC	TGGCCCTCGT	TTAGACGTCC	GTCAGGGAAG	
ACTGCAACCT	GGCTTTGGAT	GTGGTCAGCT	TTTCCAAAGG	ACACTTCAAG	CTTCTGAAAG	480
TGACGTTGGA	CCGAAACCTA	CACCAGTCGA	AAAGGTTTCC	TGTGAAGTTC	GAAGACTTGC	
TGAAAGTGGA	GGTGAGAGAC	ATTAATGACC	ATAGCCCTCA	CTTTCCAGT	GAAATAATGC	540
ACTTTCACCT	CCACTCTCTG	TAATTACTGG	TATCGGGAGT	GAAAGGGTCA	CTTTATTACG	
ATGTGGAGGT	GTCTGAAAGT	TCCTCTGTGG	GCACCAGGAT	TCCTTTAGAA	ATTGCAATAG	600
TACACCTCCA	CAGACTTTCA	AGGAGACACC	CGTGGTCTA	AGGAAATCTT	TAACGTATC	
ATGAAGATGT	TGGGTCCAAC	TCCATCCAGA	ACTTTCAGAT	CTCAAATAAT	AGCCACTTCA	660
TACTTCTACA	ACCCAGGTTG	AGGTAGGTCT	TGAAAGTCTA	GAGTTTATTA	TCGGTGAAGT	
GCATTGATGT	GCTAACCCAGA	GCAGATGGGG	TGAAATATGC	AGATTTAGTC	TTAATGAGAG	720
CGTAACTACA	CGATTGGTCT	CGTCTACCC	ACTTTATACG	TCTAAATCAG	AATTACTCTC	
AACTGGACAG	GGAAATCCAG	CCAACATACA	TAATGGAGCT	ACTAGCAATG	GATGGGGGTG	780
TTGACCTGTC	CCTTTAGGTC	GGTTGTATGT	ATTACCTCGA	TGATCGTTAC	CTACCCCCAC	
TACCATCACT	ATCTGGTACT	GCAGTGGTTA	ACATCCGAGT	CCTGGACTTT	AATGATAACA	840
ATGGTAGTGA	TAGAOCATGA	CGTCACCAAT	TGTAGGCTCA	GGACCTGAAA	TTACTATTGT	
GCCAGTGT	TGAGAGAGC	ACCATTGCTG	TGGACCTAGT	AGAGGATGCT	CCTCTGGGAT	900
CGGGTCACAA	ACTCTCTTGG	TGGTAACGAC	ACCTGGATCA	TCTCCTACGA	GGAGACCCTA	
ACCTTTTGTT	GGAGTTACAT	GCTACTGACG	ATGATGAAGG	AGTGAATGGA	GAAATGTGTT	960
TGGAAAACAA	CCTCAATGTA	CGATGACTGC	TACTACTTCC	TCACTTACCT	CTTTAACAAA	
ATGGATTGAG	CACCTTTGGCA	TCTCAAGAGG	TACGTCAGCT	ATTTAAATTT	AACTCCAGAA	1020
TACCTAAGTC	GTGAAACCGT	AGAGTTCTCC	ATGCAAGTGA	TAAATTTTAA	TTGAGGTCTT	

Figure 6A
SUBSTITUTE SHEET (RULE 26)

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CTGGCAGTGT TACTCTTGAA GGCCAAGTTG ATTTTGAGAC CAAGCAGACT TACGARTTG	1080
GACCGTCACA ATGAGAACTT CCGGTTCAAC TAAAACTCTG GTTCGTCTGA ATGCTTAAAC	
AGGTACAAGC CCAAGATTTG GGCCCCAACC CACTGACTGC TACTTGTAAG GTAAGTGTTC	1140
TOCATGTTTG GGTTCATAAC CCGGGGTTGG GTGACTGACG ATGACATTTT CATTGACAAG	
ATATACTTGA TGTAATGAT AATACCCAG CCATCACTAT TACCCCTCTG ACTACTGTAA	1200
TATATGAACT ACATTTACTA TTATGGGGTC GGTAGTGATA ATGGGGAGAC TGATGACATT	
ATGCAGGAGT TGCCATATTT CCAGAAACAG CCACAAAGGA GAACTTTATA GCTCTGATCA	1260
TACGTCTCA ACGGATATAA GGTCTTTGTC GGTGTTTCTT CTTGAAATAT CGAGACTAGT	
GCACTACTGA CAGAGCCTCT GGATCTAATG GACAAGTTCT CTGTACTCTT TATGGACATG	1320
CGTGATGACT GTCTCGGAGA CCTAGATTAC CTGTTCAAGC GACATGAGAA ATACCTGTAC	
AGCACTTTAA ACTACAGCAA GCTTATGAGG ACAGTTACAT GATAGTTACC ACCTCTACTT	1380
TCGTGAAAT TGATGTGTT CGAATACTCC TGTCATGTA CTATCAATGG TGGAGATGAA	
TAGACAGGGA AAACATAGCA GCGTACTCTT TGACAGTAGT TGCAGAAGAC CTTGGCTTCC	1440
ATCTGTCCCT TTTGTATCGT CGCATGAGAA ACTGTATCA ACGTCTTCTG GAACCGAAGG	
CCTCAATTGA GACCAAAAAG TACTACACAG TCAAGGTTAG TGATGAGAA GACAATGCAC	1500
GGAGTAACCT CTGGTTTTTC ATGATGTGTC AGTTCCAATC ACTACTCTTA CTGTTACGTG	
CTGTATTTTC TAAACCCAG TATGAAGCTT CTATTCTGGA AAATAATGCT CCAGGCTCTT	1560
GACATAAAG ATTTGGGGTC ATACTTCGAA GATAAGACCT TTTATTACGA GTCCCGAGAA	
ATATACTAC AGTGATAGCC AGAGACTCTG ATAGTGATCA AAATGGCAAA GTAAATTACA	1620
TATATTGATG TCACTATCGG TCTCTGAGAC TATCACTAGT TTTACCGTTT CATTAAATGT	
GACTTGTGGA TGCAAAAGTG ATGGGCCAGT CACTAACAAAC ATTTGTTTCT CTTGATGCGG	1680
CTGAACACCT ACGTTTTTAC TACCGGTCA GTGATGTGTT TAAACAAAGA GAACTACGCC	
ACTCTGGAGT ATTGAGAGCT GTTAGGTCTT TAGACTATGA AAACTTAAA CAACTGGATT	1740
TGAGACCTCA TAACTCTCGA CAATCCAGAA ATCTGATACT TTTTGAATTT GTTGACCTAA	
TTGAAATTGA AGCTGCAGAC AATGGGATCC CTCAACTCTC CACTCGCGTT CAACTAAATC	1800
AACITTAATC TCGACGTCTG TTACCCTAGG GAGTTGAGAG GTGAGCGCAA GTTGATTAG	
TCAGAAATAG TGATCAAAAT GATAATTGCC CTGTGATAAC TAATCCTCTT CTTAATAATG	1860
AGTCTTATCA ACTAGTTTTA CTATTAACGG GACACTATTG ATTAGGAGAA GAATTATTAC	
GCTCGGGTGA AGTTCCTGCT CCCATCAGCG CTCTCAAAA CTATTTAGTT TTCCAGCTCA	1920
CGAGCCCACT TCAAGACGAA GGGTAGTCGC GAGGAGTTTT GATAAATCAA AAGGTCGAGT	
AAGCCGAGGA TTCAGATGAA GGGCACAAC CCCAGCTGTT CTATACCATA CTGAGAGATC	1980
TTGGGCTCCT AAGTCTACTT CCGGTGTTGA GGGTCGACAA GATATGGTAT GACTCTCTAG	
CAAGCAGATT GTTTGCCATT AACAAAGAAA GTGGTGAAGT GTTCCTGAAA AAACAATTAA	2040
GTTCGTCTAA CAACGGTAA TTGTTTCTTT CACCACTCA CAAGGACTTT TTTGTTAATT	
ACTCTGACCA TTCAGAGGAC TTGAGCATAG TAGTTGCAGT GTATGACTTG GGAAGACCTT	2100
TGAGACTGGT AAGTCTCCTG AACTCGTATC ATCAACGTCA CATACTGAAC CCTTCTGGAA	
CATTATCCAC CAATGCTACA GTTAAATTCA TCTCACCGA CTCTTTTCTT TCTAACGTG	2160
GTAATAGGTG GTTACGATGT CAATTTAAGT AGGAGTGGCT GAGAAAAGGA AGATTGCAAC	

Figure 6B
SUBSTITUTE SHEET (RULE 26)

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AAGTCGTTAT	TTTGCAACCA	TCTGCAGAAG	AGCAGCAOCCA	GATCGATATG	TCATTATAT	2220
TTGAGCAATA	AAACGTTGGT	AGACGTCTTC	TOGTCGTGGT	CTAGCTATAC	AGGTAATATA	
TCATTGCAGT	GCTGGCTGGT	GGTGTGCTT	TGCTACTTTT	GGCCATCTTT	TTTGTGGCCT	2280
AGTAACGTCA	CGACOGACCA	CCAACACGAA	ACGATGAAAA	CCGGTAGAAA	AAACACCGGA	
GTACTTGTAA	AAAGAAAGCT	GGTGAATTTA	AGCAGGTACC	TGAACAACAC	GGAACATGCA	2340
CATGAACATT	TTTCTTTGCA	CCACTTAAAT	TCGTCCATGG	ACTTGTGTG	CCTTGTACGT	
ATGAAGAAGC	CCTGTTAAGC	ACCCCATCTC	CCCAGTCGGT	CTCTTCTTCT	TTGTCTCAGT	2400
TACTTCTTGC	GGACAATTGC	TGGGGTAGAG	GGGTGAGCCA	GAGAAGAAGA	AACAGAGTCA	
CTGAGTCATG	CCAACCTCTC	ATCAATACTG	AATCTGAGAA	TTGCAGCGTG	TCCTCTAACC	2460
GACTCAGTAC	GGTTGAGAGG	TAGTTATGAC	TTAGACTCTT	AACGTGCGAC	AGGAGATTGG	
AAGAGCAGCA	TCAGCAAACA	GGCATAAAGC	ACTCCATCTC	TGTACCATCT	TATCACACAT	2520
TTCTCGTCGT	AGTCGTTTGT	CCGTATTTGC	TGAGGTAGAG	ACATGGTAGA	ATAGTGTGTA	
CTGGTTGGCA	CCTGGACAAT	TGTGCAATGA	GCATAAGTGG	ACATTCTCAC	ATGGGGCACA	2580
GACCAACCGT	GGACCTGTTA	ACACGTTACT	CGTATTCAAC	TGTAAGAGTG	TACCCCGTGT	
TTAGTACAAA	GGTACAGTGG	GCAAAGGAGA	TAGTGAATTC	AATGACAGTG	ACTCTGATAC	2640
AATCATGTTT	CCATGTCACC	CGTTTCTCT	ATCACTGAAG	TTACTGTCAC	TGAGACTATG	
TAGTGGAGAA	TCAGAAAAGA	AGAGCATTGA	GCAGCCAAAG	CAGGCACAAG	CCAGTGCTCA	2700
ATCACCTCTT	AGTCCTTTCT	TCTCGTAACT	CGTCGGTTAC	GTCGGTGTTC	GGTCACGAGT	
ATACACAGAT	GAATCAGCAG	GGTTCOGACA	TGCGGATAAC	TATTTGAGCC	ACCGAATCAA	2760
TATGTGTCTA	CTTAGTCGTC	CCAAGGCTGT	ACGGCTATTG	ATAAAGTCGG	TGGCTTAGTT	
CAAGGGTCCA	GAAATGGGA	ACTGCACATT	GCAATATGAA	AAGGGCTATA	GACTGTCTTA	2820
GTCCACAGGT	CTTTTACOCCT	TGACGTGTAA	CGTTATACTT	TTCCOGATAT	CTGACAGAAT	
CTCTGTAGCT	CCTGTATATT	ACAATAOCTA	CCATGCAAGA	ATGCCTAACC	TGCACATACC	2880
GAGACATCGA	GGACATATAA	TGTTATGGAT	GGTACGTTCT	TACGGATTGG	ACGTGTATGG	
GAAOCATAAC	CTTAGAGACC	CTTATTACCA	TATCAATAAT	CCTGTTGCTA	ATCGGATGCA	2940
CTTGGTATGG	GAATCTCTGG	GAATAATGGT	ATAGTTATTA	GGACAACGAT	TAGCCTACGT	
GGCGGAATAT	GAAAGAGATT	TAGTCAACAG	AAGTGCAACG	TTATCTCCGC	AGAGATCGTC	3000
COGCTTATA	CTTTCTCTAA	ATCAGTTGTC	TTACGTTGTC	AATAGAGGCG	TCTCTAGCAG	
TAGCAGATAC	CAAGAATTCA	ATTACAGTCC	GCAGATATCA	AGACAGCTTC	ATCCTTCAGA	3060
ATCGTCTATG	GTCTTAAAGT	TAATGTCAGG	CGTCTATAGT	TCTGTGGAAG	TAGGAAGTCT	
AATTGCTACA	ACCTTTTAAAT	CATTAGGCAT	GCAAGTGAGA	ATGCACAAAG	GCAAGTGCTT	3120
TTAACGATGT	TGGAAAATTA	GTAATCOGTA	CGTTCACCTT	TACGTGTTTC	CGTTCACGAA	
TAGCATGAAA	GCTAAATATA	TGGAGTCTCC	CCTTTCCCTC	TGATGGATGG	GGGGAGACAC	3180
ATCGTACTTT	CGATTTATAT	ACCTCAGAGG	GGAAAGGGAG	ACTACCTAAC	CCCCTCTGTG	
AGGACAGTGC	ATAAATATAC	AGCTGCTTTC	TATTTGCATT	TCACTTGGGA	ATTTTTTGTT	3240
TCCTGTACAG	TATTTATATG	TCGACGAAAG	ATAAACGTAA	AGTGAACOCCT	TAAAAACAA	
TTTTTTACAT	ATTTATTTTT	CCTGAATTGA	ATGTGACATT	GTCCTGTCAC	CTAACTAGCA	3300
AAAAAATGTA	TAAATAAAAA	GGACTTAACT	TACACTGTAA	CAGGACAGTG	GATTGATCGT	

Figure 6C
SUBSTITUTE SHEET (RULE 26)

11/18

ATTAAATCCA CAGACCTACA GTCAAATATT TGAGGGCCCC TGAAACAGCA CATCAGTCAG 3360
TAATTTAGGT GTCTGGATGT CAGTTTATAA ACTCCCGGGG ACTTTGTCGT GTAGTCAGTC

GACCTAAAGT GGCCTTTTAA CTTTTCAGCAG CTCCTGGGTC TGCCCTCTGT GTTAATCAGC 3420
CTGGATTTC ACGGAAAAAT GAAATCGTC GAGGACCCAG ACGGGAGACA CAATTAGTCG

CCCTGGTCAA GTCTGAGTA GGATCATGGC GTTTTATAT GCATCTCACC TACTTTGGAC 3480
GGGACCAATT CAGGACTCAT CCTAGTACCG CAAAAATATA CGTAGAGTGG ATGAAACCTG

GTGATTYACA CATAATAGGA AACGCTTGGT TTCAGTGAAG TCTGTGTTGT ATATATTCTG 3540
CACTAAATGT GTATTATCCT TTGCGAACCA AAGTCACTTC AGACACAACA TATATAAGAC

TTATATACAC GCATTTTGTG TTTGTGTATA TATTTCAAGT CCATTCAGAT ATGTGTATAT 3600
AATATATGTG CGTAAACAC AAACACATAT ATAAAGTTCA GGTAAGTCTA TACACATATA

AGTGCAGACC TTGTAAATTA AATATTCTGA TACTTTTTCC TCAATAAATA TTTAAAT
TCACGTCTGG AACATTTAAT TTATAAGACT ATGAAAAGG AGTTATTTAT AAATTA

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MVCCGPGRML LGWAGLLVLA ALCLLQVPGA QAAACEPVRI PLCKSLPWNM TKMPNHLHHS	60
TQANAILAME QFEGLLGTHC SPDLLFPLCA MYAPICTIDF QHEPIKPCKS VCERARQGCE	120
PILIKYRHSW PESLACDELP VYDRGVCISP EAIVTADGAD FPMDSSSTGHC RGASSERCKC	180
KPVRATQKTY FRNNYNYVIR AKVKEVKMRC HDVTAVVEVK EILKASLVNI PRDTVNLYTT	240
SGCLCPPLTV NEEYVIMGYE DEERSRLLLV EGSTAEKWKD RLGKKVKRWD MKLRHLGLGK	300
TDASDSTQNO KSGRNSNPRP ARS.	

13/18

AAGCCTGGGA	CCATGGTCTG	CTGCGGCCCG	GGACGGATGC	TGCTAGGATG	GGCCGGGTTG	60
TTCGGACCCT	GGTACCAGAC	GACGCCGGGC	CCTGCCTACG	ACGATCCTAC	CCGGCCCAAC	
CTAGTCCTGG	CTGCTCTCTG	CCTGCTCCAG	GTGCCCCGAG	CTCAGGCTGC	AGCCTGTGAG	120
GATCAGGACC	GACGAGAGAC	GGACGAGGTC	CACGGGCCTC	GAGTCCGACG	TCGGACACTC	
CCTGTCCGCA	TCCCGCTGTG	CAAGTCCCTT	CCCTGGAACA	TGACCAAGAT	GCCCAACCAC	180
GGACAGGCGT	AGGGCGACAC	GTTCAGGGAA	GGGACCTTGT	ACTGGTTCTA	CGGGTTGGTG	
CTGCACCACA	GCACCCAGGC	TAACGCCATC	CTGGCCATGG	AACAGTTCGA	AGGGCTGCTG	240
GACGTGGTGT	CGTGGGTCCG	ATTGCGGTAG	GACCGGTACC	TTGTCAAGCT	TCCCAGACGAC	
GGCACCCTACT	GCAGCCCGGA	TCTTCTCTTC	TTCTCTGTGT	CAATGTACGC	ACCCATTTGC	300
CCGTGGGTGA	CGTCGGGCCT	AGAAGAGAAG	AAGGAGACAC	GTTACATGCG	TGGGTAAACG	
ACCATCGACT	TCCAGCACGA	GCCCATCAAG	CCCTGCAAGT	CTGTGTGTGA	GCGCGCCCGA	360
TGGTAGCTGA	AGGTCGTGCT	CGGGTAGTTC	GGGACGTTCA	GACACACACT	CGCGCGGGCT	
CAGGGCTGCG	AGCCCATTTCT	CATCAAGTAC	CGCCACTCGT	GGCCGGAAAG	CTTGGCCTGC	420
GTCCCGACGC	TCGGGTAAAG	GTAGTTTCATG	GCGGTGAGCA	CCGGCCTTTC	GAACCGGACG	
GACGAGCTGC	CGGTGTACGA	CCGCGGCGTG	TGCATCTCTC	CTGAGGCCAT	CGTCACCGCG	480
CTGCTCGACG	GCCACATGCT	GGCGCCGCAC	ACGTAGAGAG	GACTCCGGTA	GCAGTGGCGC	
GACGGAGCGG	ATTTTCCTAT	GGATTCAAGT	ACTGGACACT	GCAGAGGGGC	AAGCAGCGAA	540
CTGCCTCGCC	TAAAAGGATA	CCTAAGTTCA	TGACCTGTGA	CGTCTCCCCG	TTCGTGCTTT	
CGTTGCAAAT	GTAAGCCTGT	CAGAGCTACA	CAGAAGACCT	ATTTCCGGAA	CAATTACAAC	600
GCAACGTTTA	CATTCCGACA	GTCGATGT	GTCTTCTGGA	TAAAGGCCTT	GTTAATGTTG	
TATGTCATCC	GGGCTAAAGT	TAAAGAGGTA	AAGATGAAAT	GTCATGATGT	GACCGCGTTT	660
ATACAGTAGG	CCCGATTTCA	ATTTCTCCAT	TTCTACTTTA	CAGTACTACA	CTGGCGGCAA	
GTGGAAGTGA	AGGAAATTCT	AAAGGCATCA	CTGGTAAACA	TTCCAAGGGA	CACCGTCAAT	720
CACCTTCACT	TCCTTTAAGA	TTTCCGTAGT	GACCATTGTG	AAGGTTCCCT	GTGGCAGTTA	
CTTTATACCA	CCTCTGGCTG	CCTCTGTCTT	CCACTTACTG	TCAATGAGGA	ATATGTCATC	780
GAAATATGGT	GGAGACCGAC	GGAGACAGGA	GGTGAATGAC	AGTTACTCCT	TATACAGTAG	
ATGGGCTATG	AAGACGAGGA	ACGTTCCAGG	TTACTCTTGG	TAGAAGGCTC	TATAGCTGAG	840
TACCCGATAC	TTCTGCTCCT	TGCAAGGTCC	AATGAGAACC	ATCTTCCGAG	ATATCGACTC	
AAGTGGAAGG	ATCGGCTTGG	TAAGAAAGTC	AAGCGCTGGG	ATATGAAACT	CCGACACCTT	900
TTCACCTTCC	TAGCCGAACC	ATTCTTTTCAG	TTCGCGACCC	TATACTTTGA	GGCTGTGGAA	
GGACTGGGTA	AAACTGATGC	TAGCGATTCC	ACTCAGAATC	AGAAGTCTGG	CAGGAACTCT	960
CCTGACCCAT	TTTGACTACG	ATCGCTAAGG	TGAGTCTTAG	TCTTCAGACC	GTCCTTGAGA	

Figure 8A
SUBSTITUTE SHEET (RULE 26)

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AATCCCCGGC	CAGCACGCAG	CTAAATCCTG	AAATGTAAAA	GGCCACACCC	ACGGACTCCC	1020
TTAGGGGCCC	GTCGTGCGTC	GATTTAGGAC	TTTACATTTT	CCGGTGTGGG	TGCCTGAGGG	
TTCTAAGACT	GGCGCTGGTG	GACTAACAAA	GGAAAACCGC	ACAGTTGTGC	TCGTGACCGA	1080
AAGATTCTGA	CCGCGACCAC	CTGATTGTTT	CCTTTTGGCG	TGTCAACACG	AGCACTGGCT	
TTGTTTACCG	CAGACACCGC	GTGGCTACCG	AAGTTACTTC	CGGTCCCCTT	TCTCCTGCTT	1140
AACAAATGGC	GTCTGTGGCG	CACCGATGGC	TTCAATGAAG	GCCAGGGGAA	AGAGGACGAA	
CTTAATGGCG	TGGGGTTAGA	TCCTTTAATA	TGTTATATAT	TCTGTTTCAT	CAATCACGTG	1200
GAATTACCGC	ACCCCAATCT	AGGAAATTAT	ACAATATATA	AGACAAAGTA	GTTAGTGAC	
GGGACTGTTC	TTTTGCAACC	AGAATAGTAA	ATTAAATATG	TTGATGCTAA	GGTTTCTGTA	1260
CCCTGACAAG	AAAACGTTGG	TCTTATCATT	TAATTTATAC	AACTACGATT	CCAAAGACAT	
CTGGACTCCC	TGGGTTTAAT	TTGGTGTTC	GTACCCTGAT	TGAGAAATGCA	ATGTTTCATG	1320
GACCTGAGGG	ACCCAAATTA	AACCACAAGA	CATGGGACTA	ACTCTTACGT	TACAAAGTAC	
TAAAGAGAGA	ATCCTGGTCA	TATCTCAAGA	ACTAGATATT	GCTGTAAGAC	AGCCTCTGCT	1380
ATTCTCTCT	TAGGACCACT	ATAGAGTTCT	TGATCTATAA	CGACATTCTG	TCGGAGACGA	
GCTGCGCTTA	TAGTCTTG TG	TTGTATGCC	TTTGTCCATT	TCCCTCATGC	TGTGAAAGTT	1440
CGACGCGAAT	ATCAGAACAC	AAACATACGG	AAACAGGTAA	AGGGAGTACG	ACACTTTCAA	
ATACATGTTT	ATAAAGGTAG	AACGGCATTT	TGAAATCAGA	CACTGCACAA	GCAGAGTAGC	1500
TATGTACAAA	TATTTCCATC	TTGCCGTAAA	ACTTTAGTCT	GTGACGTGTT	CGTCTCATCG	
CCAACACCAG	GAAGCATTTA	TGAGGAAACG	CCACACAGCA	TGACTTATTT	TCAAGATTGG	1560
GGTTGTGGTC	CTTCGTAAAT	ACTCCTTTGC	GGTGTGTCGT	ACTGAATAAA	AGTTCTAACC	
CAGGCAGCAA	AATAAATAGT	GTTGGGAGCC	AAGAAAAGAA	TATTTTGCCT	GGTTAAGGGG	1620
GTCCGTGCTT	TTATTTATCA	CAACCTCGG	TTCTTTTCTT	ATAAAACGGA	CCAATTCCCC	
CACACTGGAA	TCAGTAGCCC	TTGAGCCATT	AACAGCAGTG	TTCTTCTGGC	AAGTTTTTGA	1680
GTGTGACCTT	AGTCATCGGG	AACTCGGTAA	TTGTGCTCAC	AAGAAGACCG	TTCAAAAAC	
TTTGTTCATA	AATGTATTCA	CGAGCATTTAG	AGATGAACTT	ATAACTAGAC	ATCTGTTGTT	1740
AAACAAGTAT	TTACATAAGT	GCTCGTAATC	TCTACTTGAA	TATTGATCTG	TAGACAACAA	
ATCTCTATAG	CTCTGCTTCC	TTCTAAATCA	AACCCATTGT	TGGATGCTCC	CTCTCCATT	1800
TAGAGATATC	GAGACGAAGG	AAGATTTAGT	TTGGGTAACA	ACCTACGAGG	GAGAGGTAAG	

Figure 8B
SUBSTITUTE SHEET (RULE 26)

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ATAAATAAAT TTGGCTTGCT GTATTGGCCA GGAAAAGAAA GTATTAAAGT ATGCATGCAT 1860
TATTTATTTA AACCGAACGA CATAACCGGT CCTTTTCTTT CATAATTTCA TACGTACGTA

GTGCACCAGG GTGTTATTTA ACAGAGGTAT GTAACCTCTAT AAAAGACTAT AATTTACAGG 1920
CACGTGGTCC CACAATAAAT TGTCTCCATA CATTGAGATA TTTTCTGATA TTAAATGTCC

ACACGGAAAT GTGCACATTT GTTTACTTTT TTTCTTCCTT TTGCTTTGGG CTTGTGATTT 1980
TGTGCCTTTA CACGTGTAAA CAAATGAAAA AAAGAAGGAA AACGAAACCC GAACACTAAA

TGGTTTTTGG TGTGTTTATG TCTGTATTTT GGGGGGTGGG TAGGTTTAAG CCATTGCACA 2040
ACCAAAAACC ACACAAATAC AGACATAAAA CCCCCACCC ATCCAAATTC GGTAACGTGT

TTCAAGTTGA ACTAGATTAG AGTAGACTAG GCTCATTGGC CTAGACATTA TGATTTGAAT 2100
AAGTTCAACT TGATCTAATC TCATCTGATC CGAGTAACCG GATCTGTAAT ACTAAACTTA

TTGTGTTGTT TAATGCTCCA TCAAGATGTC TAATAAAAGG AATATGGTTG TCAACAGAGA 2160
AACACAACAA ATTACGAGGT AGTTCTACAG ATTATTTTCC TTATACCAAC AGTTGTCTCT

CGACAACAAC AACAAA
GCTGTTGTTG TTGTTT

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MVCGSPGGML LLRAGLLALA ALCLLRVPGA RAAACEPVRI PLCKSLPWNM TKMPNHLHHS	60
TQANAILAIE QFEGLLGTHC SPDLLFFLCA MYAPICTIDF QHEPIKPCKS VCERARQGCE	120
PILIKYRHSW PENLACEELP VYDRGVCISP EAIVTADGAD FPMDSNGNC RGASSERCKC	180
KPIRATQKTY FRNNYNYVIR AKVKEIKTKC HDVTAVVEVK EILKSSLVNI PRDTVNLYTS	240
SGCLCPPLNV NEEYIIMGYE DEERSRLLLV EGSIAEKWKD RLGKKVKRWD MKLRHLGLSK	300
SDSSNSDSTQ SQKSGRNSNP RQARN.	

Figure 9
SUBSTITUTE SHEET (RULE 26)

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GGCGGAGCGG	GCCTTTTGGC	GTCCACTGCG	CGGCTGCACC	CTGCCCCATC	TGCCGGGATC	60
CCGCCTCGCC	CGGAAAACCG	CAGGTGACGC	GCCGACGTGG	GACGGGGTAG	ACGGCCCTAG	
ATGGTCTGCG	GCAGCCCCGG	AGGGATGCTG	CTGCTGCGGG	CCGGGCTGCT	TGCCCTGGCT	120
TACCAGACGC	CGTCGGGCCC	TCCCTACGAC	GACGACGCCC	GGCCCGACGA	ACGGGACCGA	
GCTCTCTGCC	TGCTCCGGGT	GCCCGGGGCT	CGGGCTGCAG	CCTGTGAGCC	CGTCCGCATC	180
CGAGAGACGG	ACGAGGCCCA	CGGGCCCCGA	GCCCGACGTC	GGACACTCGG	GCAGGCGTAG	
CCCCGTGCA	AGTCCCTGCC	CTGGAACATG	ACTAAGATGC	CCAACCACCT	GCACCACAGC	240
GGGGACACGT	TCAGGGACGG	GACCTTGTAC	TGATTCTACG	GGTTGGTGGA	CGTGGTGTCTG	
ACTCAGGCCA	ACGCCATCCT	GGCCATCGAG	CAGTTTGAAG	GTCTGCTGGG	CACCCACTGC	300
TGAGTCCGGT	TGCGGTAGGA	CCGGTAGCTC	GTCAAGCTTC	CAGACGACCC	GTGGGTGACG	
AGCCCCGATC	TGCTCTTCTT	CCTCTGTGCC	ATGTACGCGC	CCATCTGCAC	CATTGACTTC	360
TCGGGGCTAG	ACGAGAAGAA	GGAGACACGG	TACATGCGCG	GGTAGACGTG	GTAAGTGAAG	
CAGCACGAGC	CCATCAAGCC	CTGTAAGTCT	GTGTGCGAGC	GGGCCCCGCA	GGGCTGTGAG	420
GTCGTGCTCG	GGTAGTTCTG	GACATTTCAGA	CACACGCTCG	CCCGGGCCGT	CCCGACACTC	
CCCATACTCA	TCAAGTACCG	CCACTCGTGG	COGGAGAACC	TGGCCTGCGA	GGAGCTGCCA	480
GGGTATGAGT	AGTTTCATGG	GGTGAGCACC	GGCCTCTTGG	ACCGGACGCT	CCTCGACGGT	
GTGTACGACA	GGGGCGTGTG	CATCTCTCCC	GAGGCCATCG	TTACTGCGGA	CGGAGCTGAT	540
CACATGCTGT	CCCCGCACAC	GTAGAGAGGG	CTCCGGTAGC	AATGACGCCT	GCCTCGACTA	
TTTCCTATGG	ATTCTAGTAA	CGGAAACTGT	AGAGGGGCAA	GCAGTGAACG	CTGTAAATGT	600
AAAGGATACC	TAAGATCATT	GCCTTTGACA	TCTCCCCGTT	CGTCACTTGC	GACATTTACA	
AAGCCTATTA	GAGCTACACA	GAAGACCTAT	TTCCGGAACA	ATTACAACCTA	TGTCATTCCG	660
TTCCGATAAT	CTCGATGTGT	CTTCTGGATA	AAGGCCCTGT	TAATGTTGAT	ACAGTAAGCC	
GCTAAAGTTA	AAGAGATAAA	GACTAAGTGC	CATGATGTGA	CTGCAGTAGT	GGAGGTGAAG	720
CGATTTC AAT	TTCTCTATTT	CTGATTCACG	GTAATACACT	GACGTCATCA	CCTCCACTTC	
GAGATTCTAA	AGTCCTCTCT	GGTAAACATT	CCACGGGACA	CTGTCAACCT	CTATACCAGC	780
CTCTAAGATT	TCAGGAGAGA	CCATTGTGTA	GGTGCCCTGT	GACAGTTGGA	GATATGGTCG	
TCTGGCTGCC	TCTGCCCTCC	ACTTAATGTT	AATGAGGAAT	ATATCATCAT	GGGCTATGAA	840
AGACCGACGG	AGACGGGAGG	TGAATTACAA	TTACTCCTTA	TATAGTAGTA	CCCGATACTT	

Figure 10A
SUBSTITUTE SHEET (RULE 26)

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GATGAGGAAC	GTTCCAGATT	ACTCTTGGTG	GAAGGCTCTA	TAGCTGAGAA	GTGGAAGGAT	900
CTACTCCTTG	CAAGGTCTAA	TGAGAACCAC	CTTCCGAGAT	ATCGACTCTT	CACCTTCCTA	
CGACTCGGTA	AAAAAGTTAA	GCGCTGGGAT	ATGAAGCTTC	GTCATCTTGG	ACTCAGTAAA	960
GCTGAGCCAT	TTTTTCAATT	CGCGACCCTA	TACTTCGAAG	CAGTAGAACC	TGAGTCATTT	
AGTGATTCTA	GCAATAGTGA	TTCCAATCAG	AGTCAGAAAGT	CTGGCAGGAA	CTCGAACCCC	1020
TCACTAAGAT	CGTTATCACT	AAGGTGAGTC	TCAGTCTTCA	GACCGTCCTT	GAGCTTGGGG	
CGGCAAGCAC	GCAACTAAAT	CCCGAAATAC	AAAAAGTAAC	ACAGTGGACT	TCCTATTAAG	1080
GCCGTTTCGTG	CGTTGATTTA	GGGCTTTATG	TTTTTTCATTG	TGTCACCTGA	AGGATAATTC	
ACTFACTTGC	ATTGCTGGAC	TAGCAAAGGA	AAATTGCACT	ATTGCACATC	ATATTCTATT	1140
TGAATGAACG	TAACGACCTG	ATCGTTTCCT	TTTAACGTGA	TAACGTGTAG	TATAAGATAA	
GTTTACTATA	AAAATCATGT	GATAACTGAT	TATTACTTCT	GTTTCTCTTT	TGGTTTCTGC	1200
CAAATGATAT	TTTTAGTACA	CTATTGACTA	ATAATGAAGA	CAAAGAGAAA	ACCAAAGACG	
TTCTCTCTTC	TCTCAACCCC	TTTGTAATGG	TTTGGGGGCA	GACTCTTAAG	TATATTGTGA	1260
AAGAGAGAAG	AGAGTTGGGG	AAACATTACC	AAACCCCGT	CTGAGAATTC	ATATAACACT	
GTTTCTATT	TCACTAATCA	TGAGAAAAAC	TGTTCTTTTG	CAATAATAAT	AAATTAAACA	1320
CAAAAGATAA	AGTGATTAGT	ACTCTTTTTG	ACAAGAAAAC	GTTATTATTA	TTTAATTTGT	
TGCTGTTACC	AGAGCCTCTT	TGCTGAGTCT	CCAGATGTTA	ATTTACTTTC	TGCACCCCAA	1380
ACGACAATGG	TCTCGGAGAA	ACGACTCAGA	GGTCTACAAT	TAAATGAAAG	ACGTGGGGTT	
TTGGGAATGC	AATATTGGAT	GAAAAGAGAG	GTTTCTGGTA	TTCACAGAAA	GCTAGATATG	1440
AACCCTTACG	TTATAACCTA	CTTTCTCTC	CAAAGACCAT	AAGTGTCTTT	CGATCTATAC	
CCTTAAAAACA	TACTCTGCCG	ATCTAATTAC	AGCCTTATTT	TGTATGCCT	TTTGGGCATT	1500
GGAATTTTGT	ATGAGACGGC	TAGATTAATG	TCGGAATAAA	AACATACGGA	AAACCCGTAA	
CTCCTCATGC	TTAGAAAGTT	CCAAATGTTT	ATAAAGGTAA	AATGGCAGTT	TGAAGTCAAA	1560
GAGGAGTACG	AATCTTTCAA	GGTTTACAAA	TATTTCCATT	TTACCGTCAA	ACTTCAGTTT	
TGTCACATAG	GCAAAGCAAT	CAAGCACCAG	GAAGTGTTTA	TGAGGAAACA	ACACCCAAGA	1620
ACAGTGATATC	CGTTTCGTTA	GTTTCGTGGT	CTTCACAAAT	ACTCCTTTGT	TGTGGGTCT	
TGAATTATTT	TTGAGACTGT	CAGGAAGTAA	AATAAATAGG	AGCTTAAGAA	AGAACATTTT	1680
ACTTAATAAA	AACCTGTACA	GTCCTTCATT	TTATTTATCC	TGAATTCTT	TCTTGTAATA	
GCCTGATTGA	GAAGCACAAAC	TGAAACCACT	AGCCGCTGGG	GTGTTAATGG	TAGCATTCCT	1740
CGGACTAACT	CTTCGTGTTG	ACTTTGGTCA	TCGGCGACCC	CACAATTACC	ATCGTAAGAA	
CTTTTGCCAA	TACATTTGAT	TTGTTCAATG	ATATATTAAT	CAGCATTAGA	GAAATGAATT	1800
GAAAACCGTT	ATGTAAACTA	AACAAGTACT	TATATAATTA	GTCGTAATCT	CTTTACTTAA	
ATAACTAGAC	ATCTGCTGTT	ATCACCATAG	TTTTGTTTAA	TTTGCTTCCT	TTTAAATAAA	1860
TATTGATCTG	TAGACGACAA	TAGTGGTATC	AAAACAAATT	AAACGAAGGA	AAATTTATTT	
CCCATTGGTG	AAAGTCAAAA	AAAAAAAAAA	AAA			
GGGTAACCAC	TTTCAGTTTT	TTTTTTTTTT	TTT			

Figure 10B
SUBSTITUTE SHEET (RULE 26)

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US97/10942

A. CLASSIFICATION OF SUBJECT MATTER

IPC(6) : Please See Extra Sheet.

US CL : 530/300, 350; 514/2; 536/23.1

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

U.S. : 530/300, 350; 514/2; 536/23.1

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

DIALOG (MEDLINE, BIOSIS, EMBASE, WPI, USPATFULL) AUTHOR AND WORD. search terms: e.g. cerberus, xenopus

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Y, P	BOUWMEESTER et al. Cerberus is a head-inducing secreted factor expressed in the anterior endoderm of Spemann's organizer. Nature. 15 August 1996, Vol. 382, No. 6592, pages 595-601, see entire document.	1-15

☐ Further documents are listed in the continuation of Box C. ☐ See patent family annex.

* Special categories of cited documents:	
A document defining the general state of the art which is not considered to be of particular relevance	*T* later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
E earlier document published on or after the international filing date	*X* document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
L document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)	*Y* document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art
O document referring to an oral disclosure, use, exhibition or other means	*A* document member of the same patent family
P document published prior to the international filing date but later than the priority date claimed	

Date of the actual completion of the international search

29 AUGUST 1997

Date of mailing of the international search report

11 SEP 1997

Name and mailing address of the ISA/US
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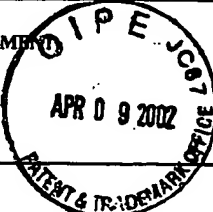
INTERNATIONAL SEARCH REPORT

International application No.
PCT/US97/10942

A. CLASSIFICATION OF SUBJECT MATTER:

IPC (6):

A01N 37/18; A61K 38/00; C07K 1/00, 2/00, 4/00, 7/00, 14/00, 16/00, 17/00; C07H 21/02, 21/04

FORM 1449* INFORMATION DISCLOSURE STATEMENT IN AN APPLICATION (Use several sheets if necessary)		Docket Number: 510015-258	Application Number: 09/903,188
		Applicant: De Robertis et al.	
		Filing Date: July 11, 2001	Group Art Unit: 1647

TECH CENTER 1600/2900

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U.S. PATENT DOCUMENTS						
EXAMINER INITIAL	DOCUMENT NO.	DATE	NAME	CLASS	SUBCLASS	FILING DATE IF APPROPRIATE
PR	5,457,048	10/10/1995	Pasquale et al.			
FOREIGN PATENT DOCUMENTS						
	DOCUMENT NO.	DATE	COUNTRY	CLASS	SUBCLASS	TRANSLATION
						YES NO
PR	94/05791	03/17/1994	PCT			
PR	94/05800	03/17/1994	PCT			
OTHER DOCUMENTS (Including Author, Title, Date, Pertinent Pages, Etc.)						
PR	Bouwmeester et al., "Cerberus is a head-inducing secreted factor expressed in the anterior endoderm of Spemann's organizer," <i>Nature</i> , 382:6592, pp. 595-601 (15 August 1996)					
	Christian et al., "Interactions between <i>Xwnt-8</i> and Spemann organizer signaling pathways generate dorsoventral pattern in the embryonic mesoderm of <i>Xenopus</i> ," <i>Genes & Development</i> , 7, pp. 13-28 (1993)					
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PR	Alberts et al., <i>Molecular Biology of the Cell</i> , January 1994, Garland Publishing, Inc., New York, NY, page(s) G-6, G-9, G-17, G-23, 1142, and 1144-1145					

EXAMINER <i>James R. Rowe</i>	DATE CONSIDERED <i>5/26/94</i>
EXAMINER: Initial if reference considered, whether or not citation is in conformance with MPEP 609; draw line through citation if not in conformance and not considered. Include copy of this form for next communication to the Applicant.	